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Dictyostelium discoideum (strain AX-3) DNA
Dictyostelium discoideum
                        M59747.1 GI:167723
                                 D.discoideum protein kinase 4 M59747
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Acylated insulin
                                                                                       Similarity
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1 (bases 1 to 87)

Haribabu, B. and Dottin, R.P.
Identification of a protein kinase multigene family of Dictyostelium discoideum: molecular cloning and expression cDNA encoding a developmentally regulated protein kinase Proc. Natl. Acad. Sci. U.S.A. 88 (4), 1115-1119 (1991)
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/product="protein kinase 4"
/protein_id="AAA33189.1"
/db_xxef="GI:167724"
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/strain="AX-3"
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Neoteleostei;

Actinopterygii; Neopterygii; Perconorpha; Pleuronectiformes;

Acanthomorpha; Acanthopterygii; Perconorpha; Pleuronectiformes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   h 0.5%; Score 25.8; DB 6; Length 81. Similarity 57.0%; Pred. No. 2.6e+05;
                                                                                                                                                                                                                                                                        Quattro,J.M. and Jones,W.J.
Direct Submission
Submitted (22-SEP-1997) Biological Sciences, University of South
Carolina, Coker Life Sciences Building, Columbia, SC 29208, USA
                                                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 96)
Quattro, J.M. and Jones, W.J.
Amplification Primers That Target Locus-Specific Introns in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AF025984 96 bp DNA linear VRT 30-OCT-: Paralichthys dentatus lactate dehydrogenase (LDHA) gene, allele
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22 c
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Candida albicans
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AX486379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Roemer,T., Jiang,B., Boone,C., Bussey,H. and Ohlsen,K.L. Gene disruption methodologies for drug target discovery patent: WO 02053728-A 3679 11-JUL-2002;
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                                                                                                                                                                                                                                                                                                                                                              Sequence 3 from Patent WO0045176
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                                                                                                                                                    Gallusser,A., Karl,J., Lill,H., Stahl,P., Krueger,K. and Borgya,A. Method of identifying n-terminal probnp Patent: WO 0045176-A 03-AUG-2000; GALLUSSER ANDREAS (DE); KARL JOHANN (DE); LILL HELMUT (DE); STAHL PETER (DE); KRUEGER KERSTIN (DE); BORGYA ANNELIESE (DE); ROCCHE DIAGNOSTICS GMBH (DE)
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Escherichia coli
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/db_xref="taxon:5476"
a 5 c 22 g 24 t
                                                                              /organism="Escherichia coli"
/db_xref="taxon:562"
19 c 21 g 20 t
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                Score 25.2; DB 1; Length 70; pred. No. 3.8e+05;
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3026 ATGTCCTGACTAAATGTGAGAGTGGTAGAGGTACAAGACCCCAGGTGGGCGACTCAAGGAT 3085
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                                                         51;
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Meagher, R.B. and Summers, A.O.
Metal resistance sequences and transgenic plants
Patent: US 5668294-A 5 16-SEP-1997;
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                                                                   Similarity
                                                                                                                                                                                                                                     Unclassified.
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21 c 25 g
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19 c 21 g
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Pred. No. 3.8e+05;
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                                                          Murphy,B.R., Collins,P.L., Schmidt,A.C., Durbin,A.P., Skiadopoulos,M.H. and Tao,T.
Use of recombinant parainfluenza viruses (pivs) as vectors to protect against infection and disease caused by piv and other human
         The Secretary of the Department of Health and Human Services (US)
                               pathogens
Patent: WO 0142445-A 31 14-JUN-2001;
                                                                                                                                         artificial sequences.
1 (bases 1 to 92)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Draft entry and computer-readable sequence for [1] kindly provided by S.H.Chang, 0.2\text{-}\text{FEB-}1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lee,C.P., Kao,M.C., French,B.A., Putney,S.D. and Chang,S.H. The rabbit muscle phosphofructokinase gene. Implications for protein structure, function, and tissue specificity J. Biol. Chem. 262 (9), 4195-4199 (1987)
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18 of 22
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KEYWORDS
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193480/c
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                                                                                                                                                source
                        3958 GAACATGTGATTAGGCTGTATTCAAAGCAGATCACCATTGCGATCAACGTCCTCC 4012
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                                                                                                                                                                                                                                                                                                                                                                                              56 GANGACGTAAGTAAGTATTAAAAACATACCCCCCCTGGCCGATCGTCCTCC 2
Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gold,L., Janjic,N. and Pagratis,N. High affinity oligonucleotide ligands to growth factors Patent: US 6207816-A 78 27-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 78 from patent US 6207816. AR140801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      h 0.5%; Score 24.8; D
Similarity 63.3%; Pred. No. 4.9e
38; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AR140801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unknown.
Unclassified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unknown.
                                                                                                                                                                                                                                                                                                 71 bp
sequence 78 from patent US 5731144.
                                                                                                                                                                           1 (bases 1 to 71)
Toothman, p.J., Ringquist, S. and Gold, L.
High affinity TGF. beta. nucleic acid ligands
Patent: US 5731144-A 78 24-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                 Unclassified.
                                                                                                                                                                                                                                                 Unknown
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                                                        Conservative
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/db_xref="taxon:32630"

/note="Reverse primer for RSV A G gene insert"

12 c 26 g 34 t
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16 c 22 g
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                                                                                                                          /organism="unknown"
16 c 22 g
                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.5%; Score 24.6; DB 6; Length 71; 65.5%; Pred. No. 5.5e+05;
                                                                   0.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches 19;
                                                      0;
                                                                     Score 24.6; DB 6; Length 71; pred. No. 5.5e+05;
                                                           Mismatches
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                                                             Indels
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DEFINITION
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S76509/c
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                                                                                                                                                                    4971 GGTTTATTTTA 4981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93 GTTTATCATACATAAGACCGCATGACTATAATACAAAGGGGGGTTCTTTTTCATTTAAC 34
                                                                                                                                           33 GTTACAATATA 23
                                                                                                                                                                                                                                                             Local
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Sequence 78 from patent US 5731424.
195107
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Toothman,P.J., Ringquist,S. and Gold,L.
High affinity TGF.beta. nucleic acid ligands and inhibitors
Patent: US 5731424-A 78 24-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              36; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 from patent US 6365352. AR203036
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                                                                                                                                                                                                                                                                                                                                                                  Process to study changes in gene expression in granulocytic cells patent: US 6365352-A 21 02-APR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Unclassified
                                                                                                                                                                                                                                                                                                                                                                                                                                                Unknown
                                                                                                                                                                                                                                                  42;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unknown
                                                 DIS8 (A) (hypervariable minisatellite MS32) [human, MACH, Genomic, 73 nt].
                          S76509.1 GI:243382
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Homo sapiens individual MACH.
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1. .71
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16 c 20 g
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                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                  Score 24.6; DB 6; Length 98; 
pred. No. 5.6e+05; 
0; Mismatches 29; Indels
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                                                                           individual
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ORIGIN
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MEDLINE
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1362 TCTGACTTCTATTTATAGACCATTTGTAGACAAAGCACTGAAGCAGATGGGGTTAAGAAA 1421
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Best Local Similarity 60.6%;
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                                                       46;
                                                                                                                                                                                                                             P-element-induced interallelic gene conversion of insertions and deletions in Drosophila melanogaster Mol. Cell. Biol. 13 (11), 7006-7018 (1993)
                                                                                                                                                                                                                           8413290
                                                                                                                                                                                                                                                                                      Johnson-Schlitz, D.M. and Engels, W.R.
                                                                                                                                                                                                                                                                                                              Unpublished (1993)
2 (bases 1 to 98)
                                                                                                                                                                                                                                                                                                                                                                                insertion sequence.

Drosophila melanogaster (individual_isolate B20.3) DNA.

Drosophila melanogaster

Drosophila melanogaster

Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;

Meoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;

Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                          Deletion white allele. Transposase-induced derivative of
                                                                                                                                                                                                                                                                                                                                                                 Engels, W.R. and Johnson_Schlitz, D.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              L20950.1 GI:304832
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Drosophila melanogaster DNA sequence, complete insertion sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GenBank staff at the National Library of Medicine created this entry [NCBI gibbsq 76509] from the original journal article. This sequence comes from Figure 2.
                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Monckton,D.G. and Jeffreys,A.J.
Minisatellite 'isoallele' discrimination in pseudohomozygotes by
Single molecule PCR and variant repeat mapping
Genomics 11 (2), 465-467 (1991)
                                                                                                                                37 a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 73)
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                                                                                                                     /db_xref="taxon:7227"
2 c 1 g
                                                                                                                                                  /organism="Drosophila melanogaster"
/isolate="B20.3"
                                                                                                                                                                                                 Location/Qualifiers
                                0.5%; Score 24.4; DB 3; Length 98; 56.1%; Pred. No. 6.4e+05; ative 0; Mismatches 36; Indels
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/db_xref="taxon:9606"
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Pred. No. 6.3e+05;
0; Mismatches 26; Indels
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                                                                                     Query Match
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                                                                                                                                                                                          JOURNAL
             4838 TTTTAATCTCCTGTTTGTTGAGTGCACTGACTGTGAAACCTTTACCTTTTTTGTTGTTGT 4897
                                                                                                                                                                                                          TITLE
                                                                                                                                                                                                                 AUTHORS
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AR007408/c
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1 TTTTTAGCTCCTGGTTTCTTAGTGCCACCTCTATGGGCTTTATTAGATTGTTGTTGTTGT 60
                                                                                                                                                                                                                                                                                                                                                                                               4529 TATTTACATAAAGACTGTGCTGAGAAGCAG 4558
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                                                                                                                                                                              Roemer, T., Jiang, B., Boone, C., Bussey, H
Gene disruption methodologies for drug
Patent: WO 02053728-A 3637 11-JUL-2002;
                                                                                                                                                                     Elitra Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                      Candida albicans
                                                                                                                                                                                                       Roemer, T., Jiang, B.,
                                                                                                                                                                                                               EUKATYOTA; Fungi, Ascomycota; Saccharomycotina; Saccharomycetes; Saccharomycetales; mitosporic Saccharomycetales; Candida.
                                                                                                                                                                                                                                                               Candida albicans.
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Acylated insulin
Patent: US 5750497-A 9 12-MAY-1998;
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12 c 13 q
                                                                                                                                 /organism="Candida albicans"
                                                                                                                                                            Location/Qualifiers
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                                              0.5%; Score 24.2; DB 6; Length 65; 62.3%; Pred. No. 7.1e+05; rative 0; Mismatches 23; Indels
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23 c 22 g
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RESULT 19
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Local Similarity 75.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 51)
Johnston,S.L. and Wettstein,P.J.
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U92173.1 GI:2894946
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Johnston, S.L. and Wettstein, P.J.
                                                                                                                                                                                                         65 bp DNA Sequence 671 from Patent WO02053728. AX483371
Roemer,T., Jiang,B., Boone,C., Bussey,H. and Ohlsen,K.L. Gene disruption methodologies for drug target discovery patent: WO 02053728-A 671 11-JUL-2002; Elitra pharmaceuticals, Inc. (US)
                                                                                                                                                                                         AX483371.1 GI:22317791
                                                                                                                                                           Candida albicans.
                                                                                                     Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; mitosporic Saccharomycetales; Candida.
                                                                                                                                           Candida albicans
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/product="T cell receptor beta chain"
/protein_id="AAC02878.1"
/protein_id="AAC02878.1"
/db_xref="GI:2894947"
/translation="YCTCSDPDRVSGNTLYF"
/translation="YCTCSDPDRVSGNTLYF"
13 c 11 g 15 t
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Pred. No. 7
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RESULT 21
AX207310
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             AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                           4855 TTGA 4858
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                                                                                                                                                                                                                                                                                                                                                                                                                                       61 TGGA 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Methods of identifying synthetic transcriptional and translational regulatory elements, and compositions relating to same regulatory elements, and compositions relating to same Patent: WO 0155371-A 35 02-AUG-2001; Patent: WO 0155371-A 35 02-AUG-2001; The Scripps Research Institute (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 35 from Patent WO0155371
AX207310
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mauro, V.P., Edelman, G.M., Chappell, G.M., Jones, F.S., Owens, G. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   artificial sequences.
                                                          Yuasa,K., Kanoh,Y., Okumura,K. and Omori,K.
Genomic organization of the human phosphodiesterase PDE11A gene.
Evolutionary relatedness with other PDEs containing GAF domains
Eur. J. Blochem. 268 (1), 168-178 (2001)
                                                                                                                                                                                                                                         Homo sapiens PDE11A gene, exon 11.
AB048411
AB048411.1 GI:15128469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 0.5%; Score 24; DB 6; Length 66; Similarity 60.9%; Pred. No. 8e+05;
                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                           AB04840S11
Yuasa, K. and Omori, K. Direct Submission
                                                                                                                                            Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                            Homo sapiens DNA.
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                                (bases 1 to 88)
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/db_xref="taxon:5476"
2 c 20 g 20 t
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/note="transcriptional regulatory element"
/note="1" t 2 g 17 t
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                                                                                                                 10 GAAGATGAAGGCGATGGAGGAGAAGATGAAGTGGATGGAGGAGAAGATGAAGTGGATGGA 69
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                               42;
                    BD010413
                                                                                                                                                                                                                                                                      Falco, S.Carl., Keeler, S.Jo. and Rice, J.Ann. Chimeric genes and methods for increasing the lysine and threonine content of the seeds of plants
Patent: US 5773691-A 63 30-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                Unclassified
                                                                                                                                                                                                                                                                                                                                                             Unknown
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Sequence 63 from patent US 5773691.
AR014621
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                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                               /organism="unknown"
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/chromosome="2"
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 Mismatches

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Pred. No. 8.1e+05;
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      PAT 31-JAN-2002
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                                    BASE COUNT
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I26754
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DEFINITION Chimeric genes and methods for increasing the lysine content of the seeds of plants Chimeric genes and methods for increasing the lysine content of the seeds of plants Chimeric genes and methods genes and methods genes and methods for increasing the lysine content of the seeds of plants Chimeric of plants.
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                                                                                                                                    nutritional value of plants
Patent: US 5559223-A 54 24-SEP-1996;
                                                                                                                                                           Synthetic storage proteins with defined structure containing programmable levels of essential amino acids for improvement of the
                                                                                                                                                                                                            1 (bases 1 to 97)
Falco, S.C., Keeler, S.J. and Rice, J.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent: JP 2001502923-A 45 06-1
EI DU PONT DE NEMOURS AND CO
OS Unidentified
PN JP 2001502923-A/45
PD 06-MAR-2001
PF 27-MAR-1998 JP 1998543284
PR 27-MAR-1997 US 08/8246;
PI SAVERIO CARL FALCO, RAYMONI
                                                                                                                                                                                                                                                                   Unclassified.
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                                                                                                                                                                                                                                                                                                                                                                        Sequence 54 from patent US 5559223.
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Falco,S.C., Iii,R.E.M. and Epelbaum,S.U.
Chimeric genes and methods for increasin
seeds of plants
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JP 2001502923-A/45
06-MAR-2001
27-MAR-1998 JP 1998543284
27-MAR-1997 US 08/8246
                                                           /organism="unknown"
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Pred. No. 8.2e+05;
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AF294354S5
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Matches 42; Conservative
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AF050514
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AF294358.1 GI:10505250
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Submitted (09-AUG-2000) Molecular Biology and Genetics, University Submitted (09-AUG-2000) Molecular Biology Submitted (09-A
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1 (bases 1 to 82)
Poloumienko,A. and Blecher,S.R.
Exon-intron structure of mammalian HPRT genes
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Human endogenous retrovirus.

Human endogenous retrovirus

viruses; Retroviridae.

viruses; Retrovirid
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Human endogenous retrovirus clone 19P reverse transcriptase gene,
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/db_xref="taxon:9913"
/chromosome="X"
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Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /sex="female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /tissue_type="liver"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /gene="HPRT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             58.3%; Pred. No. 8.2e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 23.8; DB 4; Length 82; pred. No. 9.2e+05; 0; Mismatches 22; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     72 ATGTTATACTATCCA 86
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                                                                            ocal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (25-FEB-1998) Fred Hutchinson Cancer Research Center, 1100 Fairview Ave. N. Seattle, WA 98109-1024, USA Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic Acids Res. (1998) In press 2 (bases 1 to 87) Rose, T. Schultz, E.R., Henikoff, J.G., Pietrokovski, S., McCallum, C.M., and Henikoff, S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90 bp mRNA lin
Trypanosoma brucei AnTat 1.3A C2 mRNA, 5' end.
M28495
                                                                                                                                                                                                                                                                                                              Coquelet, H., Tebabi, P., Pays, A., Steinert, M. and Pays, E. Trypanosoma brucei: enrichment by UV of intergenic trans the variable surface glycoprotein gene expression site Mol. Cell. Biol. 9 (9), 4022-4025 (1989)
                                                                                                                                                                                                                                                                                                                                                                                                                      Trypanosoma brucei cDNA to mRNA.
Trypanosoma brucei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     M28495.1 GI:161984
                                                          h 0.5%; Score 23.8; DB 3; Length 90; Similarity 55.4%; Pred. No. 9.2e+05; 46; Conservative 0; Mismatches 37; Indels
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                                                                                                                                         39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /product="reverse transcriptase"
/protein_id="AAC05569.1"
/db_xref="G1:2952455"
/translation="LMSLTICQTYVGKAIKPVREQFKKCYTIH"
13 c 12 9 28 t
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/product="reve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Human endogenous retrovirus"
/db_xref="taxon:11827"
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                                                                                                                                         /gene="AnTat 1.3A"
11 c 13 g
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                                                                                                                                                                                                                           /organism="Trypanosoma brucei"
/db_xref="taxon:5691"
                                                                                                                                                                                            /gene="AnTat 1.3A"
                                                                                                                                                                     /partial
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Pred. No. 9.2e+05;
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1420 AAGTTAATTTTAAGACTTCACAA 1442

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FEATURES
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ORGANISM
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AX326634
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                                                                                                                                                                                                                                                                                                                                            4577 TCCAAGACTGAAGACTGCACAGGTGACAAGCGTCACTTCTCCTGCTGCTGCTGTTTTGTC 4635
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                                                                                                                                                                                                                                                                                                               80 TCTAACACTTCTGTCTGCCCGGAAGACACGAGCACTCTCGCCTCCTGCTCCAGCTTGCC 22
                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
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                                Patent: WO 0192579-A 55 06-DEC-2001;
                      PE Corporation (NY) (US)
                                                      and amplification
                                                        Methods for detecting target nucleic acids using coupled ligation
                                                                                                                                                      Homo sapiens
                                                                                                                                                                                               AX326634.1 GI:18097397
                                                                                     Wenz,H.M. and Schroth,G.P.
                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                       Sequence 55 from Patent WO0192579. AX326634
                                                                                                                                                                                                                                                                                                                                                                                                 37;
                                                                                                                                                                          numan
                                                                                                                                                                                                                                                    AX326634
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10021369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       coli tumour suppressor
Curr. Biol. 9 (2), 105-108 (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (12-NOV-1998) van ES J.H., Immunology, Unieversity
Hospital Utrecht, Heidelberglaan 101 Utrecht, 3508 GA, THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           van Es, J.H.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        house mouse.
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                                                                                                                                                                                                                                                                                                                                                                                          Conservative
   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /usedin=AJ130783:apc2_CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /product="APC2 protein"
/number=2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /gene="APC2"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /gene="APC2"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                              0.5%; Score 23.8; DB 10; Length 91; 62.7%; Pred. No. 9.2e+05;
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REFERENCE
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                                                                                                                                                                                                                                                                                                                  34 GTTGCTCAGGGTGGAAGAGCTGG 12
                                                                                                                                                                                                                                                                                                                                                                       94 TGCCAATGGTGTAGTGCCCTCGGGCATAGTTATTGGCAGCATCTTCCTTGCCTGTGATAA 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4074 ATTAATCAAACTGGGAGATTTTG 4096
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 94)
Fleming,J.C., Pahl,H.L., Gonzalez,D.A., Smith,T.F. and Tenen,D.G.
Structural analysis of the CD11b gene and phylogenetic analysis of
                                                                                                                                                                                                            CD11b-leukocyte integrin alpha chain [human, Genomic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                         46;
                                                                                                                                                                                                        segment 19 of 31].
                                                                                                                       Homo sapiens.
                                                                                                                                            19 of 31
                                                                                                                                                                         S52191.1 GI:263036
                                                                                                                                                                                                                                           S52152S19
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Map location: 16.
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Quattro,J.M. and Jones,W.J.
Amplification Primers That Target Locus-Specific Introns in
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37; Conserv
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Quattro, J.M. and Jones, W.J.
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/allele="PdLDHA3"
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HOMO sapiens partial TCRBV13SZJ7S2 gene for T-cell receptor beta
Chair, junctional region, patient V DA, clone #23.
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Submitted (18-NOV-2000) May E., Internal Medicine/Rheumatology, Submitted (18-NOV-2000) May E., Internal Me
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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1\ \ (bases\ 1\ to\ 87) Beattle, K.L. Microfabricated, flowthrough porous apparatus for discrete detection of binding reactions
                                                                                                                                                                                                                                                                                                        Sequence 4 from patent US 5843767
AR062856
                                                                                                                                          Unclassified
                                                                                                                                                                             Unknown
                                                                                                                                                                                                                 Unknown.
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<1. .>
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/isolate="patient V DA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /codon_start=1
/product="I"-cell receptor beta chain"
/protein_id="CAC16917.1"
/db_xref="GI:11323071"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /gene="TCRBV13S2J2S2"
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536 AAAGAGGACAAGAAAATA 553
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               7 TTGCCACTCTAGCCTGGGTAATAGAGCAAGANTCAGTCAAAAAGAAAGAAGANGAAAGANAGA 66
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 y Match 0.5%;
Local Similarity 57.6%;
                                                                                        41;
                                                                                                                                                                                                                                                                    Cycles Denaturation Annealing Ext C 10 sec. 58 C 10 sec. 72 C 20 sec. 31 54 C 10 sec. 72 C 20 sec. 31 Gel: Acrylamide 7%, Formamide 32%, Urea 34%
                                                                                                                                                                                                                                                                                                                                                                          e-mail: sts@corona.med.utah.edu
Primer A: GCCTGGGTAATAGAGCAAG
Primer B: CAACATTTTCCCCTTACACC
                                                                                                                                                                                                                                                                                                                              Initial Denaturation: 94C 300sec
                                                                                                                                                                                                                                                                                                                                                         PCR Profile:
                                                                                                                                                                                                                                                                                                                                                             End to Label: Primer B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l (bases 1 to 88) Gerken,S.C., Matsunami,N., Plaetke,R., Albertsen,H., Ballard,L., Melis,R., Lawrence,E., Moore,M., Holik,P.R., Carlson,M., Zhao,X., Robertson,M., Bradley,P., Elsner,T., Tingey,A., Lalouel,J.-M. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted by:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genetic and physical mapping of simple sequence repeat containing sequence tagged sites from the human genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STS; PCR primer; STS sequence; microsatellite DNA; microsatellite marker; sequence tagged site; tetranucleotide repeat.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                      Conservative
                                                                                                                                                               35 a
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Eccles Institute of Human Genetics

Lake City, UT 84112
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                                                                                                                                                         /evidence=experimental
                                                                                                                                                                                            /db_xref="taxon:9606"
18. .36
                                                                                                                                                                                                               /organism="Homo sapiens"
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                                                                                                 52.6%;
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                                                                        Score 23.6; DB 11; Length 88; 
Pred. No. 1e+06; 
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AX059602
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3657 GAGGAAGGTGACCTTCAAATGGCAAAGAGGAAACAAAATTGGAGAAGGCCAGTATGGGAA 3716
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                                                              Local Similarity
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                                                                                                                                                                                                                                  M[gert,H.J., St[ndker,L. and Kreutzmann,P. Serin proteinase inhibitors Patent: WO 0078963-A 10 28-DEC-2000;
                                                                                                                                                                                                                           FORSSMANN, Wolf-Georg (DE)
                                                                                                                                                                                                                                                                                                                                unidentified
                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 10 from Patent WO0078963. AX059602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            50;
                                                                                                                                                                                                                                                                                                             unclassified.
                                                                                                                                                                                                                                                                                                                                              unidentified
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Methods for detecting target nucleic acids using coupled ligation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wenz, H.M. and Schroth, G.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                       /db_xref="taxon:32644"
/note="Mamalia"
                                                                                                                                                                         /organism="unidentified"
                                                                                                                                                                                                        Location/Qualifiers
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/Ab_xref="taxon:9606"
/Ab_xref="taxon:9606"
/note="chromosome Chr.6, 217.4 cR; GenBank number
AW675467; Unigene description/ID: splicing factor,
arginine/serine-rich 3 (SFRS3)/Hs.167460-allele of SEQ ID
NO: 25"
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AF025983
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1 (bases 1 to 100)

Quattro, J.M. and Jones, W.J.

Amplification Primers That Target Locus-Specific Introns in
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Pleuronectiformes;
Pleuronectoidei; Paralichthyidae; Paralichthys.
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Quattro, J.M. and Jones, W.J.
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                                                                                artificial sequences.
Location/Qualifiers
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Best Local Similarity 60.0%; pred. No. 1.2e+06;
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Oy 2594 AGACTGTTGACACCCTGAGAAGCATGCAGTGATAATCTTTTACTAGTTGTCATGCAGT 2653

Db 1 AGCTTGTCGACACCATGAAATGTCTGCTGCTCTGCTAGCTTTCGTGCCGT 60

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18.8 0.4 91 18.8 0.4 91 18.8 0.4 91 18.8 0.4 91 18.8 0.4 91 18.8 0.4 91	18.8 0.4 91 18.8 0.4 91 18.8 0.4 91 18.8 0.4 91	18.8 0.4 91 18.8 0.4 91 18.8 0.4 91	18.8 0.4 90 18.8 0.4 90	18.8 0.4 90 18.8 0.4 90	18.8 0.4 88 18.8 0.4 88	. 8 0.4 87 . 8 0.4 87	.8 0.4 87 .8 0.4 87	.8 0.4 87 .8 0.4 87	18.8 0.4 87 18.8 0.4 87	18.8 0.4 86 18.8 0.4 87	18.8 0.4 86	18.8 0.4 85 18.8 0.4 85	18.8 0.4 85 18.8 0.4 85	18.8 0.4 85 18.8 0.4 85	18.8 0.4 85 18.8 0.4 85	18.8 0.4 84	18.8 0.4 83	18.8 0.4 81 18.8 0.4 83 1	18.8 0.4 80	18.8 0.4 80	18.8 0.4 80 1	18.8 0.4 79 6	18.8 0.4 78 4	18.8 0.4 78 3	18.8 0.4 78 2 18.8 0.4 78 2	18.8 0.4 77 4	18.8 0.4 75 4	18.8 0.4 75 3	18.8 0.4 74 4 18.8 0.4 74 4	18.8 0.4 74 2	18.8 0.4 73 5	18.8 0.4 73 2	0.4 72 4	0.4 /1 5
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US-08-400-256-4
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Patent No. 5750497
GENERAL INFORMATION:
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                                                                 TELEPHONE: 212-867-0123
TELEPAX: 212-878-9655
INFORMATION FOR SEQ ID NO: 4
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993
     TOPOLOGY: 1; MOLECULE TYPE:
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982
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985
                                                                                            REGISTRATION NUMBER: 33,728
REFERENCE/DOCKET NUMBER: 39
TELECOMMUNICATION INFORMATION:
                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Lambiris, Elias J.
REGISTRATION NUMBER: 33,7
                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                           COMPUTER: IBM PC compation operating system: PC-DC SOFTWARE: Patentin Relactor Application Data:
                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                 APPLICANT: Andersen, Asser Siolii
APPLICANT: Markussen, Jan
TITLE OF INVENTION: ACYLATED INSULIN
                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Havelund, Svenc
APPLICANT: Halstrom, John
APPLICANT: Jonassen, Ib
                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS
                           TYPE: nucleic acid
STRANDEDNESS: sing
                                        TYPE:
                                                                                                                                                                                                                                                       STATE: N
COUNTRY:
                                                                                       TELEPHONE:
                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                        APPLICATION NUMBER: US/08/400,256 FILING DATE: 03-MAR-1995
                                              LENGTH:
                                                                                                                                                                                                                                                                        CITY: New York
                                                                                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10174-6401
                                                                                                                                                                                                                                                                 New York
                                            100 base pairs
                                                                                                                                                                                                                                                                              405 Lexington Avenue, 64th Floor
                                                                                                                                                                                                                                                 United States of America
               linear
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                                                                                                                                                                                               IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                    No. 57504970 No. 5750497disk of No. 5750497th America, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    single
                                                                                                                                                                                     Release #1.0, Version #1.25
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US-09-136-389-140
US-09-136-0-838-140
US-09-345-882-9
PCT-US93-06251-39
US-07-638-742-1
US-08-709-209-376
US-08-709-209-376
US-08-458-101-376
US-08-6458-11-376
US-08-6458-6-12
US-08-652-888-6-12
US-08-652-888-6-12
PCT-US94-06456-12
PCT-US94-06456-12
PCT-US94-06456-12
US-08-654-7378-5
US-08-654-7378-5
US-08-654-7378-5
US-08-134-346A-46
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US-08-082-842A-41
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Sequence 41, Appl
Sequence 140, App
Sequence 140, App
Sequence 140, App
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36, Appl
5, Appli
133, App
91, Appl
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1, Appli
13, Appl
43, Appl
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US-08-975-365-4
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                                                                                                                           Matches
                                                                                                                                    Best Local Similarity
                                                                                                                                                     Query Match
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 4, Application US/08975365
Patent No. 6011007
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Simi
Matches 50;
                  4529 TATTTACATAAAGACTGTGCTGAGAAGCAG 4558
                                                                              TELEPHONE: 212-867-0123
                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA.

APPLICATION NUMBER: US 08/400,256
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
37 CAATTGGAAAACTACTGTGCTTAGACGCAG 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: ACYLATED INSULIN NUMBER OF SEQUENCES: 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Andersen, A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4529 TATTTACATAAAGACTGTGCTGAGAAGCAG 4558
                                                    97 AAGTCTGACGACGCTAAGGGTATCGTTGAACAATGTTGTACTTCTATCTGTTCTTTGTAC 38
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APPLICANT:
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                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                TELEPHONE: 212-878-9655
                                                                                                                                                                                                                                                                                                                                          NAME: Lambiris, Elias J.
REGISTRATION NUMBER: 33,728
REFERENCE/DOCKET NUMBER: 391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: N
COUNTRY:
                                                                                                                                                                                                                                          TYPE:
                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/975,365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE:
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CITY: New York
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                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               405 Lexington Avenue, 64th Floor
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                                                                                                                                                                                                             linear
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                                                                                                                                                                                                                        single
                                                                                                                                0.5%; Score 26; DB 3; 55.6%; Pred. No. 7.6e+02;
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                                                                                                                  0; Mismatches
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Pred. No. 7.6e+02;
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RESULT 3
US-08-427-097-5
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                                                                                                                                                                                                                                                                                                                                                                                           В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                 Sequence 5, Application US/08878957 Patent No. 5965796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (303) 499-8089 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Meagher, Richard B.
APPLICANT: Sommers, Anne O.
TITLE OF INVENTION: Metal Resistance ?
TITLE OF INVENTION: Transgenic Plants
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Ferber, Donna M.
REGISTRATION NUMBER: 33,878
REGISTRATION NUMBER: 40
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 499-8080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL: N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "Oligonucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 99 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Ferber, Donna M.
                                                                                                                                                                                                                                                                                                                                                                                                                              3026 ATGTCCTGACTAAATGTGAGAGTGGTAGAGGTACAAGACCCAGGTGGGCGACTCAAAGGAT 3085
                                                                                                                                                                                                                                                                                                                                                      3086 TTGATTTTCTACAAGCAATTGAACCTGCCTTTAT 3119
                                                                      APPLICANT: Meagher, Richard B.
APPLICANT: Summers, Anne O.
APPLICANT: Rugh, Clayton L.
TITLE OF INVENTION: Metal Resistance Sequences and
TITLE OF INVENTION: Transgenic Plants
NUMBER OF SEQUENCES: 34
          NUMBER OF SEQUENCES: 34

CORRESPONDENCE ADDRESS:
ADDRESSEE: Greenlee, Winner and Sullivan, P.C.
STREET: 5370 Manhattan Circle, Suite 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
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                                                                                                                                                                                                                                                                                                                   62 GTGAGCATACTCAAGCAAGCCAAGTTGCCTATAT 95
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                                                                                                                                                                                                                                                                                                                                                                                           2 AAGACCCAGCTATAGGTGAAGCTGTTACTGCTGCATTCGCATGGAAGGCATTGAAGTGC 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
STREET:
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Similarity 54.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: Greenlee, Winner and Sullivan, P.C. 5370 Manhattan Circle, Suite 201
  Boulder
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Metal Resistance Sequences and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Release #1.0, Version #1.30
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 Mismatches

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; ANTI-SENSE:
US-08-878-957-5
                                CURRENT FILLNO DATE: 2000-08-16

PRIOR APPLICATION NUMBER: US 09/502,330

PRIOR APPLICATION NUMBER: US 60/502,330

PRIOR APPLICATION NUMBER: US 60/133,200

PRIOR FILLING DATE: 1999-05-07

PRIOR FILLING DATE: 1999-05-23

PRIOR FILLING DATE: 1999-03-23

PRIOR FILLING DATE: 1999-03-23

PRIOR FILLING DATE: 1999-03-23

PRIOR FILLING DATE: 1999-03-23

PRIOR FILLING DATE: 1999-03-12

PRIOR PRIOR PRIOR NUMBER: US 60/119,917

PRIOR FILLING DATE: 1399-03-12

NUMBER OF SEO ID NOS: 1304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 5
US-09-641-638-918/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 918, Application US/09641638 Patent No. 6432648 GINERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SYNTERY: COUNTRY: COUNTRY: COUNTRY: COUNTRY: COUNTRY: COUNTRY: B0303

COMPOTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: FORMAL POSSIBLE TO SYSTEM: PC-DOS/MS-DOS
MEDIUM TYPE: TRELEASE #1.0, Version #1.30
                                                                                                                                                                                                                                                 APPLICANT: Cohen, Annick
TITLE OF INVENTION: BIALLELIC MARKERS DERIVED FROM GENOMIC REGIONS CARRYING
TITLE OF INVENTION: GENES INVOLVED IN ARACHIDONIC ACID METABOLISM
FILLE REFERENCE: GENEST, 051CP1
CUBRENT APPLICATION NUMBER: US/09/641,638
CUBRENT FILING DATE: 2000-08-16
CUBRENT FILING DATE: 2000-08-16
                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
SOFTWARE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 08 FILING DATE: 21-APR-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Oligonucleotide"
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Ferber, Donna M.
REGISTRATION UNMBER: 33,878
REFERENCE/DOCKET NUMBER: 40
TELECOMMUNICATION INFORMATION.
TELEPHONE: (303) 499-8089
TELEFAX: (303) 499-8089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 800 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3026 ATGTCCTGACTAAATGTGAGAGTGGTAGAGGTACAAGACCCAGGTGGGCGACTCAAGGAT 3085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3086 TTGATTTTCTACAAGCAATTGAACCTGCCTTTAT 3119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/878,957 FILING DATE: 19-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     62 GTGAGCATACTCAAGCAAGCCAAGTTGCCTATAT 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 AAGACCCAGCTATAGGTGAAGCTGTTACTGCTGCATTCGCATGGAAGGCATTGAAGTGC 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocal Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99 base pairs
                        Patent.pm
                                                                                                                                                                                                                                                                                                                                                                                                    Bougueleret, Ly
Chumakov, Ilya
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Blumenfeld, Marta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.5%; Score 25.2; DB 2; Length 99; 54.3%; Pred. No. 1.3e+03; ive 0; Mismatches 43; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US 08/427,097
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US-08-458-423A-78/c
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                                                                 TELEFAX: (303) 793-34
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence 78, Application US/08458423A Patent No. 5731144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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                                                      SEQUENCE CHARACTERISTICS:
                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                             FILING DATE: 17-AUGUST-19 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                        FILING DATE: 8-SEPTEMBER-1993 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                 FILING DATE: 11-JUNE-1990 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/536,428
                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 536 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: HIGH AFFINITY TGFA NUCLEIC TITLE OF INVENTION: ACID LIGANDS AND INHIBITORS NUMBER OF SEQUENCES: 89
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: 12-815-391 : polymorphic base C or T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
NAME/KEY: allele
 TOPOLOGY:
            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
                                             LENGTH:
                                                                                                                                                   REGISTRATION NUMBER:
                                                                                                                                                                       NAME:
                                                                                                                                                                                            APPLICATION NUMBER: 07/931,473 FILING DATE: 17-AUGUST-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 47
                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 07/964,6; FILING DATE: 21-OCTOBER-1992
                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 07/714,131
                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             505 ACCTCAGTCTCAAAGAAAAAGACAGGGAGCAAAGAAGAAGAAA 550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  46 ACTTCAGGCACAAAGGAAAAAGRGAGGAAGTAAGGAGCCTGGGAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.5%;
Local Similarity 69.6%;
es 32; Conservation
                        nucleic acid
                                                                                                                                                                     Diane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        80111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Denver
                                        71 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8400 East Prentice Avenue,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WordPerfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STEVEN RINGQUIST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LARRY GOLD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PENELOPE J. TOOTHMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Swanson and Bratschun, L.L.C.
On Fast Prentice Avenue, Suite #200
                                                                                                                                                                  H. McClearn
         single
                                                                                                                                                                                                                                                                                                                                                                                                                                           2-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                              10-JUNE-1991
                                                                                        793-3433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MS-DOS
                                                                                                                                                                                                                                                                                                    07/964,624
                                                                                                                                                                                                                                                          08/117,991
                                                                                                                                           33,960
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    Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 24.8; DB 4; Length 47; Pred. No. 1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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US-08-458-424B-78
    Best Local Similarity 65.5%; Matches 36; Conservative
                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 7
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                                                                                                                                                                     TELEFAX: (303) 793-34 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 5731424
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 78, Application US/08458424B Patent No. 5731424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                            TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                 REGISTRATION NUMBER: 33,960
REFERENCE/DOCKET NUMBER: NE
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/931,473
FILING DATE: 17-AUGUST-1992
                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 21-OCTOE
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DALL:
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION UNBER: 07/714,131
TITME DATE: 10-JUNE-1991
                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 07/53
FILING DATE: 11-JUNE-1990
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3958 GAACATGTGATTAGGCTGTATTCAAAGCAGATCACCATTGCGATCAACGTCCTCC 4012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: HIGH AFFINITY TGFA NUCLEIC TITLE OF INVENTION: ACID LIGANDS AND INHIBITORS
                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                              TYPE:
                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 07/964,6
FILING DATE: 21-OCTOBER-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                     NAME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM CON
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            56 GAAGACGTAAGTAGTATTAAAAAACATACCCCCCTGGCCGCATCGTCCTCC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Match 0.5%;
Local Similarity 65.5%;
                                                                                                                        nucleic acid
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                                                                                                                                                                                                                                                                  Diane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SSEE: Swanson and Bratschun, L.L.C.
F: 8400 East Prentice Avenue, Suite #200
Denver
                                                                                                                                      71 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Colorado
 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LARRY GOLD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STEVEN RINGQUIST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PENELOPE J. TOOTHMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                              H. McClearn
                                                                                                         single
                                                                                                                                                                                                                                                                                                                                     8-SEPTEMBER-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Diskette, 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                     793-3433
                              0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                   07/964,624
                                                                                                                                                                                                                                                                                                                                                                                                                                             07/536,428
                                                                                                                                                                                                                                                                                                                                                       08/117,991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US/08/458,424B
0; Mismatches
           Score 24.6; DB 1;
Pred. No. 1.5e+03;
                                                                                                                                                                                                                                  NEX 34-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 24.6; DB 1; Length 71; Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                        DB 1; Length 71;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19;
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Patent No. 6207816

GENERAL INFORMATION:
APPLICANT: LARRY GOLD et al.
TITLE OF INVENTION: HIGH AFFINITY OLIGONUCLEOFIDE
TITLE OF INVENTION: LIGANDS TO GROWTH
TITLE OF INVENTION: FACTORS
                         ; TOPOLOGY: 1:
; MOLECULE TYPE:
US-08-973-124-78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                          TELEFAX: (303) 793-3433
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 536

PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/(
PTITING DATE: 30-MAY-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: MS-DOS
SOSTWARE: WORDERFECT 6.1
CURRENT APPLICATION DATA
APPLICATION NUMBER: US/08/973,124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 30-MAY-1996
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/459
FILING DATE: 02-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/47
FILING DATE: 07-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 02-JUNE-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/479,783
FILING DATE: 07-JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
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                                                                                                                                                                                REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION: TELEPHONE: (303) 793-3333
                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 71 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Englewood
STATE: Colorado
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 02-JUN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/46 FILING DATE: 05-JUNE-1995
                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: 20-MA
                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                               NAME: Barry J. Swanson REGISTRATION NUMBER: 3
                                                                               STRANDEDNESS:
                                                                                                            ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8011
                                                                                               nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Swanson & Bratschun, L.L.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Diskette, 3.5 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                     UMBER: 08/465,591
05-JUNE-1995
                                                                                                                                                                                                                                                                                   20-MARCH-1996
                                                                               single
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                                                                                                                                                                                                                                                                                                     08/618,693
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Score 24.6;
   DB 4; Length 71;
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Query Match

0.5%;

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PCT-US96-08014-78/c
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PCT-US96-08014-78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: LARRY GOLD; NEWCOOL OF THAMAN APPLICANT: PAGRATIS, PENELOPE J. TOOTHMAN FITTLE OF INVENTION: HIGH AFFINITY OLIGONUCLEOTIDE TITLE OF INVENTION: LIGANDS TO TRANSFORMING GROWTH TITLE OF INVENTION: FACTOR (TGF), PLATELET-DERIVED TITLE OF INVENTION: GROWTH FACTOR (PDGF) AND HUMAN TITLE OF INVENTION: KEROTINOCYTE GROWTH FACTOR (hKGF)
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                                                                                                                               TELEFAX: (303) 793-343
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 05-JUNE-
FRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 0
FILING DATE: 07-JUNE-
FILING DATE: 07-JUNE-
                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08
FILING DATE: 07-JUNE-1
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08
FILING DATE: 20-MARCH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM compatible OPERATING SYSTEM: MS-DOS SOFTWARE: WordPerfect 6.1 CURRENT APPLICATION NUMBER: PCT/C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/450
FILING DATE: 02-70URE-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/451
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PRIOR APPLICATION DATA:
PAPPLICATION NUMBER: 08/465,594
FILING DATE: 05-JUNE-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
PRIOR APPLICATION DATA:
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PRIOR APPLICATION NUMBER: 08/479,783
                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: (303) 793-3333
                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                             SEQUENCE CHARACTERISTICS:
                    MOLECULE TYPE:
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                                                                                                                                                                                                         REGISTRATION NUMBER: 33 REFERENCE/DOCKET NUMBER:
                                      STRANDEDNESS: single TOPOLOGY: linear
                                                                                             LENGTH:
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                                                                         nucleic acid
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                                                                                           71 base pairs
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0; Mismatches 19;
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RESULT 11
US-09-335-983-5
; Sequence 5, Application US/09335983
; Patent No. 6436628
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                    TELEAX: 201-7-1
TELEX: 13521
INFORMATION FOR SEQ ID NO: 5-
SEQUIENCE CHARACTERISTICS:
FRIGHT: 72 base pairs
                                                                                                                                                                                                                                                                             Query Match
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US-09-100-664A-5
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                                                                                                                                             3754 ATGGCCATGAA 3764
                                                                                                                                                                                     3694 ATTGGAGAAGGCCAGTATGGGAAGGTGTACACCTGCATCAGCGTCGACACCGGGGACCTG 3753
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 201-487-5800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PATENTIN Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/100,664A FILING DATE: 19-JUN-1998 CLASSIFICATION: 435 ATTORNEY/AGENT INFORMATION:
                                                                                                               61 GTGGCCATCAA 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Klauber & Jackson
                                                                                                                                                                1 ATAGGATCGGGATCGTTCGGCGACATCTACCTGGGCACCACGATCAACACTCGCGAGGAG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: A NOVEL CLOCK GENE AND METHODS OF USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: YOUNG, MICHAEL W. APPLICANT: KLOSS, BRIAN APPLICANT: BLAU, JUSTIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Jackson Esq., David A. REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Hackensack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     56 GAAGACGTAAGTAAGTATTAAAAACATACCCCCCCTGGCCGCATCGTCCTCC 2
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                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                  0.5%; Score 24.6; DB 3; Length 72; ilarity 59.2%; Pred. No. 1.6e+03; Conservative 0; Mismatches 29; Indels
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                                                                                                                                                                                                                                                                                                                                                        linear
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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EARLIER FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 66
SOUTWARE: PatentIn Ver. 2.0
SEQ ID NO 21
LENGTH: 98
TYPE: DNA
ORGANISM: Homo sapiens
US-09-506-729-21
         B
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                                                                                                                                                                                     Query Match 0.5%; Score 24.6; DB 4; Length 98; Best Local Similarity 59.2%; Pred. No. 1.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/506,729
CURRENT FILING DATE: 2000-02-18
EARLIER APPLICATION NUMBER: PCT/US98/17284
EARLIER FILING DATE: 1998-08-21
EARLIER APPLICATION NUMBER: 60/056,844
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21, Application US/09506729
Patent No. 6365352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Yerramilli, Subrahmanyam V.
APPLICANY: Prashar, Yatindra
APPLICANY: Prashar, Yatindra
APPLICANY: Newberger, Peter
APPLICANT: Goguen, Jon
APPLICANT: Weissman, Sherman M.
TITLE OF INVENTION: A PROCESS TO STUDY CHANGES IN GENE EXPRESSION IN
TITLE OF INVENTION: GRANULOCYTIC CELLS
FILE REFERENCE: 44921-5016-US
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LENGTH: 72
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                                        4971 GGTTTATTTTA 4981
                                                                                               4911 GTTTGTAATGCAAAAGGCTGATTACTGAAATTTAAGAAAAAAGGTTCTTTTTTCAATAAAT 4970
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: A NOVEL CLOCK GENE AND METHODS OF USE THEREOF FILE REPERENCE: 600-1-221N (1997-335,983)
CURRENT APPLICATION NUMBER: US/09/335,983
CURRENT FILING DATE: 1999-06-18
EARLIER APPLICATION NUMBER: 09/100,664
EARLIER FILING DATE: 1998-06-19
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Young, Michael W
APPLICANT: Kloss, Brian
APPLICANT: Blau, Justin
APPLICANT: Price, Jeffrey
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ORGANISM: Drosophila melanogaster
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33 GTTACAATATA 23
                                                                       93 GTTTATCATACATAAGACCGCATGACTATAATACAAAGGGGGGGTTCTTTTTTCATTTAAC 34
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 Mismatches

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US-08-400-256-9/c

RESULT 13

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; MOLECULE TYPE: US-08-400-256-9
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                                                                                                                                                                                                                                                                                                US-08-975-365-9/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                      Sequence 9, Application US/08975365
Patent No. 6011007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
SELENGTH: 100 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Markussen, Jan
TITLE OF INVENTION: ACYLATED INSULIN
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS: 0. 57504970 No. 5750497disk of No. 5750497th America, Inc.
ADDRESSEE: No. 57504970 No. 5750497disk of No. 5750497th America, Inc.
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MEDIUM TYPE: Floppy disk
COMPUTER: BIM PC compatible
COMPUTER: FORM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 39
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-867-0123
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STREET: New York
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                                                                                                                                                                                                                                                                                                                                                                                                    4529 TATTTACATAAAGACTGTGCTGAGAAGCAG 4558
                                                                                                                                     APPLICANT: Halstrom, John
APPLICANT: Jonassen, Ib
APPLICANT: Andersen, Asser Sloth
APPLICANT: Markussen. Tan
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                                                                   TITLE OF INVENTION: ACYLATED INSULIN
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSE: No. 60110070 No. 6011007disk of No. 6011007th America, Inc.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/400,256 FILING DATE: 03-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                    37 CAATTGGAAAACTACTGTGGTTAGACGCAG 8
                                                                                                                                                                                                                                                                                                                                                                                                                                            97 AAGTCTGACGACGCTAAGGGTATCGTTGAACAATGTTGTACTTCTATCTGTTCTTTGTAC 38
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match 0.5%; Score 24.4; DB 1; Length 100; Local Similarity 54.4%; Pred. No. 2.1e+03; Nes 49; Conservative 0; Mismatches 41; Indels
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COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New York
               New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Halstrom, John
Jonassen, Ib
Andersen, Asser Sloth
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                                                  405 Lexington Avenue, 64th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
United States of America
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US-08-975-365-9
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TELEPHONE: 212-867-0123
TELEPAX: 212-889-9655
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 100 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 15
US-08-182-175A-54
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Sequence 54. Application US/08182175A
Patent No. 555923
GENERAL INFORMATION:
APPLICANT: Saverio Carl Falco
APPLICANT: Sharon J. Keeler
APPLICANT: Janet A. Rice
TITLE OF INVENTION: Synthetic Storage Proteins with Defined Structure Containin
NUMBER OF SEQUENCES: 113
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: PSTEM: PC-TOS/MS-DOS

OPERATING SYSTEM: PC-TOS/MS-DOS

SOFTWARE: Patentin Release #1.0.
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/975,365
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FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
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                                                                                                                                                                OPERATING SYSTEM: Macintosh System, 6.0 SOFTWARE: Microsoft Word, 4.0 CURRENT APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/743,
FILING DATE: 9 August 1991
ATTORNEY/AGENT INFORMATION:
NAME: Linda Axamethy Floyd
                                                                                                                                                                                                                                                ZIP: 19898

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk
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REGISTRATION NUMBER: 33
REFERENCE/DOCKET NUMBER:
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ZIP: 19898
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                                                                                                     CLASSIFICATION:
                                                                                                                           FILING DATE:
                                                                                                                                               APPLICATION NUMBER: US/08/182,175A
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                                                                                                                                                                                                                                                                                                                                       wilmington
: Delaware
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                                                                                                                                                                                                                                                                                                                    USA
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                                                                      07/743,006
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COCATION: 2..88
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                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
         CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: BARBARA C. SIEGELL
                                                                                    COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MICROSOFT WORD VERSION 2.0C
CURRENT APPLICATION DATA:
                                                                                                                                                     COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: E. I. DU
ADDRESSEE: AND COMPA.
                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1120 GAGTATGAGGGTGATGACACAGAAGGAGATTAAAGGAGTTGGAAAGTAGTACGGATGAG 1179
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REGISTRATION NUMBER:
                                                            FILING DATE:
                                                                   APPLICATION NUMBER:
                                                                                                                                              MEDIUM TYPE:
                                                                                                                                                                                                                   CITY: WILMINGTON
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CELL TYPE: DH5 alpha
IMMEDIATE SOURCE:
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LENGTH: 97 base pairs
                                                                                                                                                                                          COUNTRY:
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TELEX: 835420
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AND COMPANY
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                                                                                                                                            FLOPPY DISK
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                                                                                                                                                                                                                                                                                                                   AND THREONINE CONTENT
                                                                                                                                                                                                                                                                                            107
                                                                   US/08/474,633A
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Pred. No. 2.7e+03;
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US-08-823-771-63
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Patent No. 6459019
GENERAL INFORMATION:
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TELEX: 835420
INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
FRIGH: 97 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local Similarity
       COMPUTER: IDM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MICROSOFT WORD VERSION 2.0C
CURRENT APPLICATION NUMBER: US/08/823,771
FILING DATE: 24-Mar-1997
CLASSIFICATION JATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/474,633
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LOCATION: 2.88
OTHER INFORMATION:
OTHER INFORMATION:
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STRAIN: E. CC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10 GAAGATGAAGGCGATGGAGGAGAAGATGAAGTGGATGGAGGAGAAGATGAAGTGGATGGA 69
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OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CELL TYPE:
                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: CHIMERIC GENES AND METHODS FOR INCREASING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLONE:
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                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: E. I. DU PONT DE NEMOURS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
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                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   42;
FILING DATE: <Unknown>
                                                                                                                                                                                                             CITY: WILMINGTON STATE: DELAWARE
                                                                                                                                                                                             COUNTRY: U.S.A.
                                                                                                                                                                                                                                      STREET: 1007 MARKET STREET
                                                                                                                                                                                                                                                        ADDRESSEE: E. I. DU PONT DE NEMOURS
AND COMPANY
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"5.11.11.5"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               storage protein
/product= "protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /function= "synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /gene= "ssp"
                                                                                                                                                                                                                                                                                                              AND THREONINE CONTENT
                                                                                                                                                                                                                                                                                                                                INCREASING THE LYSINE
                                                                                                                                                                                                                                                                                                        107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 24; DB 1;
Pred. No. 2.7e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 97;
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; SEQUENCE DESCRIPTION: SEQ ID NO: 63: US-08-823-771-63
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                                                                                                                                                                                                                                                                                                                                                                                                                                   PCT-US92-06412-54
                                                                                                                                                                                                                                                                                             Sequence 54, Application PC/TUS9206412
GENERAL INFORMATION:
APPLICANT: Saverio Carl Falco
APPLICANT: Sharon J. Keeler
APPLICANT: Janet A. Rice
TITLE OF INVENTION: Synthetic Storage Proteins with Defined Structure Containing
NUMBER OF SEQUENCES: 113
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 63:
SEQUENCE CHARACTERISTICS:
LENGTH: 97 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1120 GAGTATGAGGGTGATGACACAGAAGGAGATTAAAGGAGTTGGAAAGTAGTACGGATGAG 1179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1180 AGTGAAGAAGAA 1191
                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/743,006
FILING DATE: 9 August 1991
                                                                                                 SOFTWARE: Microsoft Word, 4.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10 GAAGATGAAGGCGATGGAGGAGAAGATGAAGTGGAGGAGGAAGATGAAGTGGATGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     70 AGAGAAGATGAA 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                          MEDIUM TYPE: Floppy Disk
COMPUTER: Macintosh
OPERATING SYSTEM: Macintosh System, 6.0
SOFTWARE: Microsoft Word 1 1
                                                                                                                                                                                                                                                    STREET: 1007 Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: 302-992-4931
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                   STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN: E. COLI
CELL TYPE: DH5 alpha
IMMEDIATE SOURCE:
                                                                  APPLICATION NUMBER: POFILING DATE: 19920807
                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.5%;
Similarity 58.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: BARBARA C. SIEGELL
REGISTRATION NUMBER: 30,684
REFERENCE/DOCKET NUMBER: BB-1037-C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: 2..88
OTHER INFORMATION:
                                                                                                                                                                                                       19898
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     storage protein
/product= "prote
/gene= "ssp"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VAME/KEY:
                                                                                                                                                                                                                                   Delaware
                                                                                                                                                                                                                                                                    E. I. du Pont de Nemours and Company
1007 Market Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                     USA
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2..88
                                                                                       PCT/US92/06412
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; Pred. No. 2.7e+03;
0; Mismatches 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /function= "synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 97;
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PCT-US92-06412-54
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TELEX: 885420

INFORMATION FOR SEQ ID NO: 5

SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
1 PNCTH: 97 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 19
US-08-672-158A-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 8, Application US/08672158A Patent No. 5770371
                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Linda Axamethy Flo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (302) 992-4929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IMMEDIATE SOURCE:
CLONE: 92-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORIGINAL SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1120 GAGTATGAGGGTGATGACACAGAAGGAGAATTAAAGGAGTTGGAAAGTAGTACGGATGAG 1179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Linda Axamethy Floyd
REGISTRATION NUMBER: 33,692
REFERENCE/DOCKET NUMBER: BB
                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 57703710 No. 5770371disk of No. 5770371th America, Inc.
                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Sheryl Thompson
TITLE OF INVENTION: Modification of Cryptic Splice Sites In
                                                                         CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Agris Dr., Cheryl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
                   NAME: Agris Dr., Cheryl H.
REGISTRATION NUMBER: 34,086
REGISTRATION NUMBER: 488
REFERENCE/DOCKET NUMBER: 487
TELEPHONE: 212-867-0123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10 GAAGATGAAGGCGATGGAGGAGAAGATGAAGTGGATGGAGGAGAAGATGAAGTGGATGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CELL TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            70 AGAGAAGATGAA 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                      STREET:
                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                  APPLICATION NUMBER: FILING DATE: 27-JU
TELEFAX: 212-878-9655
                 TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                         10174-6401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E. coli
PE: DH5 alpha
                                                                                                                                                                                                                                                                                                          New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (302) 992-4929
(302) 892-7949
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.5%; Score 24; DB 5; illarity 58.3%; Pred. No. 2.7e+03; Conservative 0; Mismatches 3
                                                                                                                                                                                                                                                                                                                                        405 Lexington Avenue, 64th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                         USA
                                                                                                                                                                                                                                                                                                                          York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA (genomic)
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                                                                                                                      27-JUN-1996
N: 435
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/product= "protein"
/gene= "ssp"
/standard_name= "5.11.11.5"
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                                                                                                                                                        US/08/672,158A
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                                                       4855.000-US
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US-08-631-751A-4
                                                                                                                   Matches
                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4, Application US/08631751A Patent No. 5843767
                                                                                                                                                                                                                                                                                                TELEFAX: (202)639-6604
INFORMATION FOR SEQ ID NO: 4:
                          3861 GTATTT 3866
                                                                 3801 CAAGGAAACTGCAGACGAATTGAAAATATTCGAAGGCATCAAACACCCCCAATCTGGTTCG 3860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                     MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                          NAME: Sanzo, Michael A.
REGISTRATION NUMBER: 36,912
REFERENCE/DOCKET NUMBER: HAN
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)639-6500
                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS
                                         83 CATTCAAATYCCTGAAGTAYTCATTATAGTCAAGGGCATATCCWACAACAAACTTGTCTG 24
                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23 GAATTT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: 11
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: 11-Apr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                           ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Vinson & Elkins
STREET: 1455 Pennsylvania Avenue, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH:
                                                                                                              38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31;
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                                                                                                                                                                                                                                                                     87 base pairs
                                                                                                             Conservative
                                                                                                                                                                                hamster, human and mouse
                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Beattie,
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                                                                                                                                                                                                           DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                             11-April-1996
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                                                                                                                         0.5%;
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72.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Microfabricated, Flowthrough Porous
Apparatus for Discrete Detection of Binding Reactions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kenneth L
                                                                                                                                                                                                                                                                                                                                                                                                                                         US/08/631,751A
                                                                                                    Score 23.6; DB 2; Length 87;
Pred. No. 3.3e+03;
2; Mismatches 26; Indels
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                                                                                                                                                                                                                                                                                                                                                        HARC0001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Version #1.25
                                                                                                     0;
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                                                                                                  Gaps
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                                                                                                  0;
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US-09-641-638-916/c
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                                                                                                            GENERAL INFORMATION:
                                                 APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                           Sequence 916, Application US/09641638 Patent No. 6432648
TITLE OF INVENTION: BIALLELIC MARKERS DERIVED FROM GENOMIC REGIONS CARRYING TITLE OF INVENTION: GENES INVOLVED IN ARACHIDONIC ACID METABOLISM
                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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Patent No. 5298400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                              2594 AGACTGTTGACACCCTGAGAAGCATGCAGGTGGATAATCTTTTACTAGTTGTCATGCAGT 2653
                                                                                                                                                                                                                                                          2654 CTGCG 2658
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid;
DESCRIPTION: Synthetic DNA oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 16
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
APPLICANT:
                                                                                                                                                                                                                          61 TGGTG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                          1 AGCTTGTCGACACCATGAAATGTCTGCTGCTCGCTCTCGGTCTGCTAGCTTTCGTGCCGT 60
                                                                                                                                                                                                                                                                                                                                                              y Match 0.5%;
Local Similarity 60.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US
FILING DATE: 19910506
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ZIP: 22313-0299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Alexandria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                 39; Conservative
                                                            Bougueleret, Lydie
Chumakov, Ilya
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEIC ACID
                                              Cohen, Annick
                                                                                            Blumenfeld, Marta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       899149
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    77 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                CDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WHITFELD, Peter L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (703)836-9300
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                                                                                                                                                                                                                                                                                                                                             Mismatches
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CURRENT APPLICATION NUMBER: US/09/641,638
CURRENT FILING DATE: 2000-08-16
PRIOR APPLICATION NUMBER: US 09/502,330
PRIOR FILING DATE: 2000-02-11
PRIOR APPLICATION NUMBER: US 60/133,200
PRIOR FILING DATE: 1999-05-07
PRIOR APPLICATION NUMBER: US 09/275,267
PRIOR APPLICATION NUMBER: US 09/275,267
PRIOR FILING DATE: 1999-03-23
PRIOR FILING DATE: 1999-03-23
PRIOR PILING DATE: 1999-03-21
PRIOR PILING DATE: 1999-02-12
NUMBER OF SEQ ID NOS: 1304
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LOCATION: 24
LOCATION: 12-815-383 : polymorphic base A or G
US-09-641-638-916
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SOFTWARE: Pau
; SEQ ID NO 916
FENGTH: 47
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                                                                              ; TOPOLOGY: un;
; MOLECULE TYPE:
US-08-643-704A-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                          REGISTRATION NUMBER: 38.4
REFERENCE/DOCKET NUMBER:
INFORMATION FOR SEQ ID NO: 11
SEQUENCE CHARACTERISTICS:
LENGTH: 60 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: MICHAELS, Mark
APPLICANT: MANN, MICHAEL
TITLE OF INVENTION: MUTANTS OF THE GREEN FLUORESCENT PROTEIN
TITLE OF INVENTION: HAVING IMPROVED FLUORESCENCE AT 37 DEGRE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: CRANDALL, Craig A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Thousand Oaks
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AUUKESSEE
                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                            FILING DATE
                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
Match 0.5%; Score 23; DB 3; Length 60; Local Similarity 68.1%; Pred. No. 4e+03; es 32; Conservative 0; Mismatches 15; Indels
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1840 DeHavilland Drive
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                                                                                                                            unknown
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                                                                                                                                                                                                                                                                  38,416
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                  0,
                    Gaps
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Sequence 9, Application US/08246583; Sequence 9, Application US/08246583; Patent No. 5750394; GENERAL INFORMATION: APPLICANT: Palese, Peter Applicant: Palese, Peter Applicant: OF INVENTION: LIBERTIFICATION TITLE OF INVENTION: COMPOUNDS THAT TITLE OF INVENTION: AND VIRAL PROTECTION AND VIRAL PROTECTION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 24
US-09-344-888A-8
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CURREMY APPLICATION NUMBER: US/09/344,888A
CURREMY APPLICATION NUMBER: 199-06-25
PRIOR APPLICATION NUMBER: EP99113156.8
PRIOR APPLICATION NUMBER: EP98119078.8
PRIOR FILING DATE: 1998-07-15
PRIOR APPLICATION NUMBER: EP98119078.8
PRIOR FILING DATE: 1998-10-09
NUMBER OF SEO ID NOS: 24
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; GENERAL INFORMATION:
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US-08-246-583-9
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LENGTH: 94
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Patent No. 6291245
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Kopetzki, Erhard
APPLICANT: Schantz, Christian
TITLE OF INVENTION: New Host-Vector System
ETITE DEFENSEMENT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                           ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
              CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                           SOPTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
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Local Similarity 60.3%; Pred. No. 5.1e+03;
Nes 38; Conservative 0; Mismatches 25
                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ZIP: 10036-2711
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                                                          APPLICATION NUMBER: US/OFFILING DATE: 20-MAY-1994
                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
NAME: Coruzzi, Laura A
                                                                                                                                                                                                                                                              NEW YORK
                                                                                                                                                                                                                                                                                                 E: PENNIE & EDMONDS
1155 AVENUE OF THE AMERICAS
                                                                                                                                                                                                                                                                                                                                                                                 IDENTIFICATION AND USE OF ANTIVIRAL COMPOUNDS THAT INHIBIT INTERACTION OF HOST CELL PROTEINS AND VIRAL PROTEINS REQUIRED FOR VIRAL REPLICATION
                                                                                US/08/246,583
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US-08-484-322-5
                                                                                        TELEPHONE: (212) 758-4800
TELERAX: (212) 751-6849
TELEX: 421792
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 96 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 26
US-08-484-322-5
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                            ORIGINAL SOURCE:
ORGANISM: hom
                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 20:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
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TITLE OF INVENTION:
TITLE OF INVENTION:
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TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
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             INDIVIDUAL ISOLATE:
                                                          TOPOLOGY:
                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  515 CAAAGAAAAAAGACAGGGAGCAAAGAGGACAAGAAAATACGT 556
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LENGTH: 76 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 69:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                               REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: June 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                 SOFTWARE:
                                                                                                                                                                                                                                                                                                                                               COMPUTER:
                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE:
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Local Similarity 71.4%;
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DEDNESS: double
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66141 PENNIE
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                                                                  single
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                                                                                                                                                                                                                                                                                                                                                       FLOPPY DISK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AMINO ACID SEQUENCES OF HYPERVARIABLE REGION 1 OF THE ENVELOPE 2 GENE OF ISOLATES OF HEPATITIS C VIRUS AND THE USE OF REAGENTS DERIVED FROM THESE HYPERVARIABLE SEQUENCES IN DIAGNOSTIC METHODS AND VACCINES
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                                                                                                                                                                                          ; ANTI-SENSE: US-07-679-052A-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-07-679-052A-10/c
                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Db
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                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 27
                                                                                                                                                                                                                                                                                            TELEX: 899149
TRIEX: 899149
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 77 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                    2598 TGTTGACACCCTGAGAAGCATGCAGGTGGATAATCTTTTACTAGTTGTCATGCAGTCTGC 2657
                               2658 G 2658
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid;
DESCRIPTION: Synthetic DNA oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: (703)836-9300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 1000
CITY: Alexandria
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: RENUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
17 G 17
                                                  77 TGTCGACACCATGAAATGTCTGCTGCTCGCTCTCGGTCTGCTAGCTTTCGTGCCGTTGGT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4201 AAAGGAGAGGGCCA 4214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4141 GAAGTGAACAGCACCCTGGGGACAGCAGCATACATGGCACCTGAAGTCATCACTCGTGGC 4200
                                                                                                                                                                                                                                                             LENGTH: 77 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 1:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                 TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: BENT, Stephen A. REGISTRATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 ACGCGGGGCGCCCA 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GAAACCTACACCACCGGGGGGGGCTGCTGGTCAGACCGCGTCTGGATTCACCAGTCTTTTC 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                            37;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22313-0299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0, Application US/07679052A 5298400
                                                                                                                          Conservative
                                                                                                                                                                                                         YES
                                                                                                                                                                                                                                                                                                                                                              (703)683-4109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1800 Diagonal Road, Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RICHARDSON, Michael A. BUNN, Clive L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WHITFELD, Peter L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Foley & Lardner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19910506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Floppy disk
                                                                                                                   0.5%; Score 22.6; DB 1;
60.7%; Pred. No. 5.9e+03;
ative 0; Mismatches 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.5%;
56.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RECOMBINANT PRODUCT
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                                                                                                                                                                                                                                                                                                                                                                                                                            29,768
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                 Length 77;
                                                                                                                     Indels
                                                                                                                0; Gaps
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RESULT 28

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US-08-123-702-21/c
; Sequence 21, Application US/08123702
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NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31.284
REFERENCE/DOCKET NUMBER: TSII
TELECOMMUNICATION INFORMATION:
TELLEPHONE: (404)-873-8795
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 90 base pairs
TYPE: NUMBER: 1000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 44; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Wadsworth
APPLICANT: Snyder, I
APPLICANT: Reddy, V¢
APPLICANT: Wei, Char
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION UNBER: US/08/123,702
PILING DATE: 17-SEPT-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Containing a Genomic DNA Insert of the KI and OX-2 Regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic Center
STREET: 1201 West Peachtree Street
CITY: Atlanta
                                                                                          APPLICANT: JEFFREYS, Alec J.
APPLICANT: ARMOUR, John
TITLE OF INVENTION: SIMPLE TANDEM REPEATS
NUMBER OF SEQUENCES: 125
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                315 ATCTTTGCCAAAAAAAATCAATTCCAGATGTGGATCTCAATAAGCCTTACCTCAGCCTTGG 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                          375 CTGTAGCAATGCTAAGCTTC 394
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: doub
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                             82 ATCTAAACCAAACAAAACCATCTCTTTGGTGAGTAACAACAATCACAACAGCAGCAGCAG 23
                                                                                                                                                                                                                                                                                                                         22 CAGCAGCCACCATTTACTTC 3
                                   STREET: 1100 New
CITY: Washington
    COUNTRY:
                                                         ADDRESSEE: CUSHMAN DARBY & CUSHMAN, L.L.P. STREET: 1100 New York Avenue, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30309-3450
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                                                                                                                                                                                                          1, Application US/08332766A 5843647
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Reddy, Vermuri, B.
Wei, Chamer
                   ₽.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wadsworth, Samuel
U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A cDNA Genomic Hybrid Sequence Encoding APP770
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.4%; Score 22.4; DB 1; Length 90; 55.0%; Pred. No. 7.4e+03; ative 0; Mismatches 36; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
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; TOPOLOGY: linear; MOLECULE TYPE: DNA (genomic) US-08-332-766A-41
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US-08-182-175A-48
Sequence 48, Application US/08182175A
Patent No. 5559223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERNCE, DOCKET NUMBER: 2172

REFERNCE, DOCKET NUMBER: 2172

TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 861-3000
TELEFAX: (202) 822-9944

TELEX: 6714627 CUSH
TELEX: 6714627 CUSH
INFORMATION FOR SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
LENGTH: 95 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Saverio Carl Falco
APPLICANT: Sharon J. Keeler
APPLICANT: Sharon J. Keeler
APPLICANT: Janet A. Rice
TITLE OF INVENTION: Synthetic Storage Proteins with Defined Structure Containin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9326052.9
FILING DATE: 21-DEC-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPETATING SYSTEM: PC-DOS/MS-DOS
SOFTWARDS. PRI-LIFE
SOFTWARDS.
                                                                                                          SOFTWARE: Microsoft Word, 4.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER OF THE PROPERTY APPLICATION NUMBER OF T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           139 AAAGAAAAAATGAATGCACCAAATCAGCCTCCACATAAAGACACTGGAAAAAACAGTGGAG 198
                                                                                                                                                                                                      ZIP: 19898
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/743,006
FILING DATE: 9 August 1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 199 AATGTGGAAGAATACAGCTA 218
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/332,766A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: BIRD,
                                                                                                                                                                                                                                                                                      STREET: 100
CITY: Wilmington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 GAGGTTGAAAAAAACAACTA 80
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es 44; Conservative
                                                                                           APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20005-3918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleic acid
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                                                                                                                                                                                                                                                                      USA
                                                                                                                                                                                        E: Floppy Disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-1994
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Pred. No. 7.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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US-08-474-633A-57
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                                                                                                                                                                                                                                                                                                                                                                                                Patent No. 5773691
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
ATTORNEY/AGENT INFORMATION:
NAME: BARBARA C. SIEGELL
                                                                          OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MICROSOFT WORD VERSION 2.0C
CURRENT APPLICATION DATA:
                                                                                                                                                 COMPUTER READABLE FORM:
                                                                                                                                                                                            STREET: 100,
CITY: WILMINGTON
                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: E. APPLICANT: COM
                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1120 GAGTATGAGGGTGATGACACAGAAGGAGAATTAAAGGAGTTGGAAAGTAGTACGGATGAG 1179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1180 AGTGAAGAAGAA 1191
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                                                                                                                                                                                                                                       ADDRESSEE:
ADDRESSEE:
                                                                                         OPERATING SYSTEM:
SOFTWARE: MICROSC
                                 CLASSIFICATION: 800
                                                 FILING DATE:
                                                       APPLICATION NUMBER: US/08/474,633A
                                                                                                                         COMPUTER:
                                                                                                                                 MEDIUM TYPE: FLOPPY DISK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: CDS
LOCATION: 2..88
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IMMEDIATE SOURCE:
CLONE: 86-H23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: 1
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 97 base pairs
TYPE: nucleic acid
                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: BB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (302) 992-4929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10 GAAGATGAAGGCGATGGAGGAGAAGCTGAAGAAGATGGAGGAGAAGCTGAAGAAGATGGA 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       70 AGAGAAGATGAA 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN: E. coli
CELL TYPE: DH5 alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
TOPOLOGY: 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER:
                                                                                                                                                                   19898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         41;
                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/08474633A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Linda Axamethy Floyd RATION NUMBER: 33,692
                                                                                                                                                                                                                     1007 MARKET STREET
                                                                                                                                                                               U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           835420
                                                                                                                                                                                                                                                                                                                                                                       COMPANY
                                                                                                                  IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                       E. I. DU PONT DE NEMOURS
AND COMPANY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (302) 892-7949
                                                                                                                                                                                                                                                                                                                                                                                   I. DU PONT DE NEMOURS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.4%; Score 22.4; DB 1; Length 97; 56.9%; Pred. No. 7.7e+03;
                                                                                                                                                                                                                                                                                        INCREASING THE LYSINE AND THREONINE CONTENT OF THE SEEDS OF PLANTS
                                                                                                                                                                                                                                                                                                                                       METHODS FOR INCREASING
                                                                                                                                                                                                                                                                                                                                                        CHIMERIC GENES AND
                                                                                                                                                                                                                                                                                 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /function= "synthetic storage protein
/product= "protein"
/gene= "ssp"
/standard_name= "5.8.8.5"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31;
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US-08-823-771-57
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                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 57, Application US/08823771
Patent NO. 6459019
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 302-773-0164
TELEX: 835420
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1180 AGTGAAGAAGAA 1191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1120 GAGTATGAGGGTGATGACACAGAAGGAGGAGTTAAAGGAGTTGGAAAGTAGTACGGATGAG 1179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: CDS
LOCATION: 2..88
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: 302-992-4931
TELEFAX: 302-773-0164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            70 AGAGAAGATGAA 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10 GAAGATGAAGGCGATGGAGGAGAAGCTGAAGAAGATGGAGGAGAAGCTGAAGAAGATGGA 69
                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MICROSOFT WORD VERSION 2.0C
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match 0.4%; Local Similarity 56.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                 APPLICATION NUMBER: US/08/823,771
FILING DATE: 24-Mar-1997
CLASSIFICATION: CUNKNOWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN: E. CELL TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLONE:
                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: E. I. I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: CHIMERIC GENES AND
                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER:
APPLICATION NUMBER: 08/474,633
                                                                                                                                                                                              COUNTRY: U.S.A.
                                                                                                                                                                                                                STATE: DELAWARE
                                                                                                                                                                                                                            CITY: WILMINGTON
                                                                                                                                                                                                                                            STREET: 1007 MARKET STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    86-H23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E. coli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           97 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                  19898
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DH5 alpha
                                                                                                                                                                                                                                                                                                                                                                                             COMPANY
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                                                                                                                                                                                                                                                                                                                                                                                                         I. DU PONT DE NEMOURS AND
                                                                                                                                                                                                                                                             AND COMPANY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /standard_name=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           storage protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /gene= "ssp"
/stand---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /function= "synthetic
                                                                                                                                                                                                                                                                                                                           AND THREONINE CONTENT
                                                                                                                                                                                                                                                                                                                                                           METHODS FOR
                                                                                                                                                                                                                                                                                                                                            INCREASING THE LYSINE
                                                                                                                                                                                                                                                                           DU PONT DE NEMOURS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30,684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 22.4; DB 1; Length 97; Pred. No. 7.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BB-1037-C
                                                                                                                                                                                                                                                                                                                                                           INCREASING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31; Indels
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SEQUENCE DESCRIPTION: SEQ ID NO: 57: US-08-823-771-57
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PCT-US92-06412-48
; Sequence 48, Application PC/TUS9206412
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1120 GAGTATGAGGGTGATGACACAGAAGGAGAATTAAAGGAGTTGGAAAGTAGTACGGATGAG 1179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1180 AGTGAAGAAGAA 1191
                                                                                                                                                                                                                                                                                                                           APPLICANT: Saverio Carl Falco
APPLICANT: Sharon J. Keeler
APPLICANT: Janet A. Rice
TITLE OF INVENTION: Synthetic Storage Proteins with Defined Structure Containing
NUMBER OF SEQUENCES: 113
                                         COMPUTER: Macintosh System, 6.0 OPERATING SYSTEM: Macintosh System, 6.0 SOFTWARE: Microsoft Word, 4.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: PCT/US92/06412 FILING DATE: 19920807
                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS
               CLASSIFICATION: 530
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10 GAAGATGAAGGCGATGGAGGAGGAGGCTGAAGAAGATGGAGGAGGAAGCTGAAGAAGATGGA 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       70 AGAGAAGATGAA 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match 0.4%; Score 22.4; DB 4; Length 97; Local Similarity 56.9%; Pred. No. 7.7e+03;
                                                                                                                                                                                                                                                CITY: Wilmington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 835420

TELEX: 835420

RMATION FOR SEQ ID NO: 57:
SEQUENCE CHARACTERISTICS:
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TELEPHONE: 302-992-4931
TELEFAX: 302-773-0164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) ORIGINAL SOURCE:
                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                               ADDRESSEE:
APPLICATION NUMBER: 07/743,006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IMMEDIATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     41; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: BARBARA C. SIEGELL REGISTRATION NUMBER: 30,684 REFERENCE/DOCKET NUMBER: BB-1037-C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  storage protein
/product= "protein"
/gene= "ssp"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: CDS
LOCATION: 2.88
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN: E. coli
CELL TYPE: DH5 alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX:
                                                                                                                                                                                                         19898
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 97 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLONE: 86-H23
                                                                                                                                                                                                                                         Delaware

    E.I. du Pont de Nemours and Company
1007 Market Street

                                                                                                                                                                                                                          USA
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                                                                                                                                                                    Floppy Disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches 31; Indels
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TELEFAX: (304,
TELEX: 835420
INFORMATION FOR SEQ ID NO: 48:
SEQUENCE CHARACTERISTICS:
LENGTH: 97 base pairs
TYPE: NUCLEIC ACID
TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCT-US92-06412-48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 34
US-08-465-591A-75
                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 5837834
Patent No. 5837834
PAURICAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 33,692
REFERENCE/DOCKET NUMBER: BB-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (302) 992-4929
TELEFAX: (302) 892-7949
TELEFAX: 0362.7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 9 August 1991 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1120 GAGTATGAGGGTGATGACACAGAAGGAGGATTAAAGGAGTTGGAAAGTAGTACGGATGAG 1179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1180 AGTGAAGAAGAA 1191
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OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: HIGH AFFINITY hKGF NUCLEIC TITLE OF INVENTION: ACID LIGANDS AND INHIBITORS NUMBER OF SEQUENCES: 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Linda Axamethy Floyd REGISTRATION NUMBER: 33,692
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: NIKOS PAGRATIS
APPLICANT: LARRY GOLD
                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5
MEDIUM TYPE: IBM compatible
COMPUTER: IBM compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 6.0
                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                  FILING DATE: 5-June 1995
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/714,131
FILING DATE: 10-JUNE-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN: E. coli
CELL TYPE: DH5 alpha
                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10 GAAGATGAAGGCGATGGAGGAGAAGCTGAAGAAGATGGAGGAGAAGCTGAAGAAGATGGA 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLONE: 86-H23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         70 AGAGAAGATGAA 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                      STREET: 8400
CITY: Denver
                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 75, Application US/08465591A
o. 5837834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    41; Conservative
                                                                                                                                                                                                                                      8011
                                                                                                                                                                                                                                                                     Colorado
                                                                                                                                                                                                                                                                                                     8400 East Prentice Avenue, Suite #200
                                                                                                                                                                                                                                                       USA
                                                                                                                                                                                                                                                                                                                         Swanson and Bratschun, L.L.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    . 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA (genomic)
                                                                                                                                                                                                 Diskette, 3.5 inch, 1.44 MB storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.4%; Score 22.4; DB 5; Length 97; 56.9%; Pred. No. 7.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /function= "synthetic storage protein /product= "protein" /gene= "ssp" ______
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /standard_name= "5.8.8.5"
                                                                                                              US/08/465,591A
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 Mismatches

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US-08-465-594A-75
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                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION UMBER: 07/714,131
FILING DATE: 10-JUNE-1991
PRIOR APPLICATION NUMBER: 07/536,428
FILING DATE: 11-JUNE-1990
PRIOR APPLICATION DATA:
**POLICATION DATA: 07/664 624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 75, Patent No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                APPLICATION NUMBER: 07/964,624
FILING DATE: 21-OCTOBER-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/117,991
FILING DATE: 8-SEPTEMBER-1993
                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (303) 793-3433
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: HIGH AFFINITY hKGF NUCLEIC TITLE OF INVENTION: ACID LIGANDS AND INHIBITORS NUMBER OF SEQUENCES: 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: NIKOS PAGRATIS
APPLICANT: LARRY GOLD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: NETELECOMMUNICATION INFORMATION: TELEPHONE: (303) 793-3333
                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/465,594A FILING DATE: 5-June 1995
                                                                                                                                                                                                                                                                                                    SOFTWARE:
                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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NAME: Diane H. McClearn
REGISTRATION NUMBER: 33,5
                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                       COMPUTER:
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 8-SEPTEMBER-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 07/964,6; FILING DATE: 21-OCTOBER-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: FILING DATE: 11-JUN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                    80111
Barry J. Swanson
                                                                                                                                                                                                                                                                                                                                                                                                                                 Denver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5, Application US/08465594A 5846713
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8400 East Prentice Avenue, Suite #200
                                                                                                                                                                                                                                                                                                                                                                                                USA
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                                                                                                                                                                                                                                                                                                                                IBM compatible
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                               8-SEPTEMBER-1993
                                                                                                                                                                                                                                                                                                                                       Diskette, 3.5 inch, 1.44 MB storage
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SER: NEX 39-2
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RESULT 36
US-08-973-124-260
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US-08-465-594A-75
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Best Local Similarity
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                                                                      APPLICATION NUMBER:
FILING DATE: 07-JUNE
PRIOR APPLICATION DATA:
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                            PRIOR APPLICATION DATA:
                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                   APPLICATION NUMBER: 08/458,424
FILING DATE: 02-JUNE-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/465,594
                                                                                                                                                                        PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                FILING DATE: 02-JUNE-1995
PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
PRIOR TOWNSON NUMBER: 08/458,423
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                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 536
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
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TITLE OF INVENTION: LIGANDS TO GROWTH
TITLE OF INVENTION: FACTORS
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 FILING DATE:
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           APPLICATION NUMBER:
                                           FILING DATE:
                                                                                                                                       APPLICATION NUMBER: 08/465,591
FILING DATE: 05-JUNE-1995
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                                                           APPLICATION NUMBER:
                                                                                                                                                                                           FILING DATE:
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REFERENCE/DOCKET NUMBER: NE
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
TELEFAX: (303) 793-3433
                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 30-MAY
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                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
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CITY: E
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LENGTH: 71 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LARRY GOLD et al. NETRONICLEOTIDE VENTION: HIGH AFFINITY OLIGONUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                                                                                                                                             WordPerfect 6.1
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                                                                                         07-JUNE-1995
                                          07-JUNE-1995
                                                                                                                                                                                    05-JUNE-1995
20-MARCH-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Diskette, 3.5 inch, 1.44 Mb storage
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                                                       08/479,783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.4%; Score 22.2; DB 2; 56.9%; Pred. No. 7.3e+03;
                                                                                                       08/479,725
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PCT-US96-08014-260
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GENERAL INFORMATION:
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TELEPAX: (303) 793-3433
INFORMATION FOR SEO ID NO: 256
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
LENGTH: 71 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
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TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: LARRY GOLD; NEBOJSA JANJIC; STEVE KINCOURS
APPLICANT: PAGRATIS; PENBLOPE J. TOOTHAN
APPLICANT: PAGRATIS; PENBLOPE J. TOOTHAN
TITLE OF INVENTION: HIGH AFFINITY OLIGONUCLEOTIDE
TITLE OF INVENTION: EACTOR (TGF), PLATELET-DERLYED
TITLE OF INVENTION: GROWTH FACTOR (PDGF) AND HUMAN
TITLE OF INVENTION: KERCTINOCYTE GROWTH FACTOR (REGF)
                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US
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MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb storage
COMPUTER: IBM compatible
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                                                                                                                                                                                                                              PRIOR APPLICATION NATA:
APPLICATION NUMBER: 08/458,423
FILLING DATE: 02-JUNE-1995
PRIOR APPLICATION NATA:
APPLICATION NATA:
PRIOR APPLICATION NATA:
PRIOR APPLICATION NATA:
08/458,424
FILING DATE: 02-JUNE-1995
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                                                                                          APPLICATION NUMBER: 08/465,591
FILING DATE: 05-JUNE-1995
PRIOR APPLICATION DATA:
                                                                                                                                                    APPLICATION NUMBER: 08/46 FILING DATE: 05-JUNE-1995 PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/479,783
FILING DATE: 07-JUNE-1995
                                                                                                                                                                                                                PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
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                                                       APPLICATION NUMBER: FILING DATE: 07-JUI
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8400 E. Prentice Avenue, Suite 200
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                                                          07-JUNE-1995
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                                                                              08/479,725
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FATCH NO. 5837456
GENERAL INCORMATION:
APPLICANT: GOLD ET AL
FITLE OF INVENTION: (
TITLE OF INVENTION: (
TITLE OF INVENTION: (
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US-08-488-402A-99/c
US-08-488-402A-99/c
; Sequence 99, Application US/08488402A
; Sequence 5837456
; Sequence 100 5837456
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Best Local :
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TELEPAX: (303) 793-43333
INFORMATION FOR SEQ ID NO: 26(SEQUENCE CHARACTERISTICS:
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TELEPHONE: (303) 793-3333
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 8400
CITY: Englewood
CTATE: Colorado
                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 08/177,99
FILING DATE: 8-SEPTEMBER-1993
ATTORNEY_AGENT INFORMATION:
NAME: Barry J. Swanson
                                                                                                                                               PRIOR DATE: 11-JUNE-1990
PRIOR APPLICATION DATA: 07/714,131
APPLICATION NUMBER: 07/714,131
FILING DATE: 10-JUNE-1991
PRIOR APPLICATION DATA: 07/931,473
APPLICATION NUMBER: 07/931,473
APPLICATION DATE: 17-AUGUST-1992
                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/964,6:
APPLICATION NUMBER: 07/968
ETILING DATE: 21-0CTOBER-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/536,428
FILING DATE: 11-JUNE-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 71 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17 GACUAUCUGGCUCGAUCCAAUCACCGGAGCCCACCGGGCCAGACGACUGGC 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31 GAGAATCITTATGGTACCTCTCCCCCCAGCACACCTCGACAGATGAAACGC 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: All pyrimidines are 2'-fluoro (2'F) modified
                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08, FILING DATE: 07 JUNE 1995 CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%;
Match
Local Similarity 56.9%;
Les 29; Conservative
                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM:
SOFTWARE: WORDPE
                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GOLD ET AL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                                      IBM pc compatible SYSTEM: MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Swanson & Bratschun, L.L.C.
00 E. Prentice Avenue, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-MARCH-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Diskette, 3 1/2 diskette, 1.44 MG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HORMONES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIGH AFFINITY OLIGONUCLEOTIDE LIGANDS TO CHORIONIC GONADOTROPIN HORMONE AND RELATED GLYCOPROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08/618,693
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                                                                                                                                                                                                                                                                                                                                                               US/08/488,402A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 22.2; DB 5; Length 71; pred. No. 7.3e+03; 4; Mismatches 18; Indels ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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US-08-484-552A-99/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 99, Application US/08484552A Patent No. 5849890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                    CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/536,428
FILING DATE: 11-JUNE-1990
PRIOR APPLICATION DATA:
                                                                                             FILING DATE: 21-OCTOBER-1992 PRIOR APPLICATION DATA:
                                                                                                                                           PRIOR APPLICATION DATA:
                                                                                                                                                                            FILING DATE: 10-JUNE-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/931,473
                                                                                                                                                                                                                                                                                                                        SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/
FILING DATE: 07 JUNE 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (303) 793-3433
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1280 TTGAGAGGCTCGAATCTGAGGATGATTCTTCTTGGCTGGGGAGCACCAGACT 1330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
REFERENCE/DOCKET NUMBER: NEX36-2
                  NAME: Barry J. Swanson REGISTRATION NUMBER: 3
                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
CITY: F
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REFERENCE,DOCKET NUMBER: NEX36-1
TELECOMMUNICATION INFORMATION:
                                                                  FILING DATE:
                                                                                APPLICATION NUMBER:
                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                            APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM pc compatible OPERATING SYSTEM: MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity es 33; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           85 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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DAVIS, KEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JAYASENA, SUMEDHA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Swanson & Bratschun, L.L.C.
                                                               8-SEPTEMBER-1993
                                                                                                                                                           17-AUGUST-1992
                                                                                                                                                                                                                                                                                                                                                                                                                  Diskette, 3 1/2 diskette, 1.44 MG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GONADOTROPIN HORMONE AND RELATED GLYCOPROTEIN HORMONES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIGH AFFINITY OLIGONUCLEOTIDE LIGANDS TO CHORIONIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            793-3333
                                                                                 08/177,991
                                                                                                                              07/964,624
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               33,215
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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PCT-US96-09472-99
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                                                                                                                                              TELEFAX: (303) 793-34
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                FEATURE:
                                                              FEATURE:
                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 85 base pairs
                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: (303) 793-3333
                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/48
FILING DATE: 07 JUNE 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (303) 793-3433
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA: 08/488,402
                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1280 TTGAGAGGCTCGAATCTGAGGATGATTCTCTTGGCTGGGGAGCACCAGACT 1330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PEATURE:
OTHER INFORMATION: All U's are 2'-F modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
         OTHER INFORMATION: All U's are 2'-F modified
                                           OTHER INFORMATION:
                                                                         TOPOLOGY: linear
                                                                                        STRANDEDNESS: sing
                                                                                                                                                                                                                                             NAME:
                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
O7 JUNE 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                           NAME: Barry J. Swanson REGISTRATION NUMBER: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 8400 5.
CITY: Englewood
                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM pc
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8400 E. Prentice Avenue, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        85 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                           WordPerfect 6.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Swanson & Bratschun, L.L.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Diskette, 3 1/2 diskette, 1.44 MG
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                                                                                      single
                                                                                                                                                                 793-3433
                                        All C's are 2'-F modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIGH AFFINITY OLIGONUCLEOTIDE LIGARDS TO CHORIONIC GONADOTROPIN HORMONE AND RELATED GLYCOPROTEIN HORMONES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.48;
64.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                            MS-DOS
                                                                                                                                                                                                                                                                                     08/484,552
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                                                                                                                                                   99:
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Pred. No. 8.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 85;
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Search completed: November 11, 2002, 10:47:56 Job time: 134 secs

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Scoring table:
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Perfect score:
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Maximum Match 100%
Listing first 1000 summaries
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Maximum DB seq length: 100
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 pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
                                                                                                                                                 Score
 26
25
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  IDENTITY_NUC Gapop 10.0 ,
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4990
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                                                                                                                                            Query
Match
                                                                                                                                                                                                                                                                                                                                                                                                   320260 seqs, 177392727 residues
                                                                                                                                                                                                              . (cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*
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	529 18.8 0.4 91 1 530 18.8 0.4 91 1	527 18.8 0.4 91 1	526 18 8 0 4 91 1	524 18.8 0.4 91 1	523 18.8 0.4 90 1	521 18.8 0.4 89 1 522 18.8 0.4 60 1	520 18.8 0.4 89 1	519 18.8 0.4 89 1	518 18.8 0.4 88 1	517 18.8 0.4 88 1	18.8 0.4 88 1	18.8 0.4 88 1	18.8 0.4 88 1	18.8 0.4 88 1	18.8 0.4 87 1	510 18.8 0.4 87 1	509 18.8 0.4 87 1	18 8 0 4 87 1	507 18 8 0.4 86 1	506 18.8 0.4 86 1	504 18.8 0.4 85	100 100 0.4 65	TO 2 10 0 0 4 05	502 18 8 0 4 62 .	501 18.8 0.4 82	500 18.8 0.4 82	18.8 0.4 81	498 18.8 0.4 81	100 10.8 0.4 80	197 19 8 0 4 80	18 8 0 4 00	495 18 8 0 4 90	494 18 8 0 / 80	493 18.8 0.4 80	492 18.8 0.4 80	18.8 0.4 80	18.8 0.4 77	18.8 0.4 77	18.8 0.4 76	18.8 0.4 75	16.6 0.4 /4	486 18 B 0 4 74	485 18 8 0 4 72	484 18.8 0 4 72	18.8 0.4 71	18.8 0.4 69	18.8 0.4 69	18.8 0.4 67	18.8 0.4 67	18.8 0.4 66	18.8 0.4 65	18.8 0.4 64	18 8 0 4 60	10.0 0.4 54	10 0 0 4 4/	18.8 0.4 45	18.8 0.4 37	10.0 0.4 32	18 8 0.4 32	100	19 0.4 100	19 0.4 100	19 0.4 100	10 0.4 99	19 0.4	19 0.4 20	19 0.4 9/	19 0.4 9/	19 0.4 95	19 0 1 06	
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	529 18.8 0.4 91 530 18.8 0.4 91	527 18.8 0.4 91 1	526 18 8 0 4 91 1	524 18.8 0.4 91 1	523 18.8 0.4 90 1	521 18.8 0.4 89 1 522 18.8 0.4 60 1	520 18.8 0.4 89 1	519 18.8 0.4 89 1	518 18.8 0.4 88 1	517 18.8 0.4 88 1	18.8 0.4 88 1	18.8 0.4 88 1	18.8 0.4 88 1	18.8 0.4 88 1	18.8 0.4 87 1	510 18.8 0.4 87 1	509 18.8 0.4 87 1	18 8 0 4 87 1	507 18 8 0.4 86 1	506 18.8 0.4 86 1	504 18.8 0.4 85	100 100 0.4 65	TO 2 10 0 0 4 05	502 18 8 0 4 62 .	501 18.8 0.4 82	500 18.8 0.4 82	18.8 0.4 81	498 18.8 0.4 81	100 10.8 0.4 80	197 19 8 0 4 80	18 8 0 4 00	495 18 8 0 4 90	494 18 8 0 / 80	493 18.8 0.4 80	492 18.8 0.4 80	18.8 0.4 80	18.8 0.4 77	18.8 0.4 77	18.8 0.4 76	18.8 0.4 75	16.6 0.4 /4	486 18 B 0 4 74	485 18 8 0 4 72	484 18.8 0 4 72	18.8 0.4 71	18.8 0.4 69	18.8 0.4 69	18.8 0.4 67	18.8 0.4 67	18.8 0.4 66	18.8 0.4 65	18.8 0.4 64	18 8 0 4 60	10.0 0.4 54	10 0 0 4 4/	18.8 0.4 45	18.8 0.4 37	10.0 0.4 32	18 8 0.4 32	100	19 0.4 100	19 0.4 100	19 0.4 100	10 0.4 99	19 0.4	19 0.4 20	19 0.4 9/	19 0.4 9/	19 0.4 95	19 0 1 06	
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APPLICANT: Kamigaki, Laura Y. (Ito)
APPLICANT: Sherman, Braddley K.
TITLE OF INVENTIO: POLYNICLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN SEEDLING FILE REFERENCE: PL-0012-1 CON 09/293,876
CURRENT APPLICATION NUMBER: US/09/923,876
CURRENT FILING DATE: 2001-08-06
PRIOR APPLICATION NUMBER: 09/298,329
PRIOR APPLICATION NUMBER: 60/085,331
PRIOR APPLICATION NUMBER: 60/085,331
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PRIOR FILING DATE: 1998-05-05
NUMBER OF SEQ ID NOS: 6332
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                                 3570 GAGGTACCGAGAAATGAGGAGAAAGAATATCATTGGTCAAGTTTGTGATACGCCTAAGTC 3629
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                                                                                  Score 26.2; DB 10;
Pred. No. 7.8e+02;
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US-09-882-434A-14

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US-09-951-401-35

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                                                                      38;
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Sequence 5162, Ap
Sequence 1837, Ap
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Sequence 59, Appl
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Sequence 112, Appl
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Sequence 9612, Ap
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Sequence 10893, A
Sequence 1032, Ap
Sequence 1078, Ap
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Sequence 14, Appl
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                                                                 Gaps
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31702, A
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR PRIOR FILING DATE: 2000-08-03
PRIOR PRIOR PRIOR NUMBER: GB 24263.6
PRIOR PRIOR PRIOR NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR PRIOR DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PILING DATE: 2001-01-30
PRIOR PRIOR PRIOR DATE: 2001-01-30
PRIOR PRIOR PRIOR DATE: 2001-01-30
PRIOR PRIOR PRIOR DATE: 2001-01-30
PRIOR PRIOR DATE: 2001-01-30
PRIOR PRIOR DATE: 2001-01-30
PRIOR RELING DATE: 2001-01-30
PRIOR PRIOR PRIOR DATE: 2001-01-30
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; NAME/KEY: misc_feature
; LOCATION: 3, 4
; OTHER INFORMATION: n = A,T,C or G
US-09-919-580-772
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/919,580
CORRENT FILING DATE: 2001-07-30
NUMBER OF SEQ ID NOS: 934
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 772
LENGTH: 81
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US-09-919-580-772/c
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                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Acomica-x-1
CURRENT FILLING DATE: 2001-05-23
CURRENT FILLING DATE: 2001-05-23
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APPLICANT: Xu, Jiangchun
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.552
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
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APPLICATION NUMBER: PCT/US01/00669
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Pred. No. 9.4e+02;
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OTHER INFORMATION: MAP TO AC007684.2

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.1

OTHER INFORMATION: EST_HUMAN HIT: A1273296.1, EVALUE 3.00e-44

OTHER INFORMATION: MIT: 414506376, EVALUE 2.00e-44

OTHER INFORMATION: SWISSPROT HIT: P41892, EVALUE 2.40e-02

US-09-864-761-31768
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
PRIOR FILING DATE: 2000-06-30
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR PELICATION NUMBER: PCT/US01/00663
PRIOR PELICATION NUMBER: PCT/US01/00665
PRIOR PILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
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                                                                                                                                                                           APPLICANT: COLLINS, Peter L.
APPLICANT: Collins, Peter L.
APPLICANT: Schmidt, Alexander C.
APPLICANT: Schmidt, Alexander C.
APPLICANT: Schmidt, Anna P.
APPLICANT: Skiadopoulos, Mario H.
APPLICANT: Skiadopoulos, Mario H.
APPLICANT: Tao. Tao
TITLE OF INVENTION: USE OF RECOMBINANT PARAINFLUENZA VIRUSES (PIVS) AS
TITLE OF INVENTION: VECTORS TO PROTECT AGAINST INFECTION AND DISEASE CAUSED
TITLE OF INVENTION: BY PIV AND OTHER HUMAN PATHOGENS
TITLE REFERENCE: 15280-404100US
TITLE REFERENCE: 15280-40410US
CURRENT FILING DATE: 2000-12-08
BRIOR APPLICATION NUMBER: 05/170,195
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEO, ID NOS: 62
COMMENT OF SECOND NOS: 62
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                                                                                                   SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO 31

LENGTH: 92
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OTHER INFORMATION: Description of Artificial Sequence: Reverse primer
                            FEATURE:
                                                 ORGANISM: Artificial Sequence
                                                                                    TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    67 AAGAAATTATTATGAAAGACTGTAAACACCCAAATATTGTTGCTTATTTTGGAAGCT 8
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similarity 61.5%;
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Pred. No.
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; OTHER INFORMATION: for RSV A G gene insert US-09-733-692A-31
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CURRENT APPLICATION NUMBER: US/09/864,761

CURRENT FILING DATE: 2001-05-23

PRIOR APPLICATION NUMBER: US 60/180,312

PRIOR FILING DATE: 2000-02-04

PRIOR FILING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR FILING DATE: 2000-08-03

PRIOR APPLICATION NUMBER: US 09/632,366

PRIOR APPLICATION NUMBER: US 09/632,366

PRIOR APPLICATION NUMBER: US 60/283,39

PRIOR APPLICATION NUMBER: US 60/286,359

PRIOR APPLICATION NUMBER: US 60/286,359

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR PILING DATE: 2000-09-27

PRIOR FILING DATE: 2000-09-27

PRIOR FILING DATE: 2000-09-27

PRIOR PILING DATE: 2000-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: PCT/US01/00664

PRIOR APPLICATION NUMBER: PCT/US01/00669

PRIOR APPLICATION NUMBER: PCT/US01/00669

PRIOR FILLING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00665

PRIOR APPLICATION NUMBER: PCT/US01/00668

PRIOR APPLICATION NUMBER: PCT/US01/00668

PRIOR APPLICATION NUMBER: PCT/US01/00663

PRIOR APPLICATION NUMBER: PCT/US01/00663

PRIOR APPLICATION NUMBER: PCT/US01/00663

PRIOR APPLICATION NUMBER: PCT/US01/00663

PRIOR APPLICATION NUMBER: PCT/US01/00663
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                                                                                                                                                   SOFTWARE: Anno
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PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
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PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 49117
FEATURE:

OTHER INFORMATION: MAP TO AC009305.1

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 0.78

OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 1

OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 0.8
                                                                                                                                   ORGANISM: Homo sapiens
                                                                                                                                                                                         LENGTH:
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                                                                                                                                                                                                                                                Annomax Sequence Listing Engine vers. 1.1
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Pred. No. 26
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US-09-864-761-28769/c
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                                  SOFTWARE:
                                                 NUMBER OF SEQ ID NOS: 49117
                                                                   PRIOR FILING DATE: 2001-01-29
                                                                                                                                                       PRIOR
                                                                                                                                                                                PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
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LENGTH: 80
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APPLICANT: Rank, David R.
APPLICANT: Hanzel, David R.
APPLICANT: Chen, Wensheng
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OTHER INFORMATION: EST HUMAN HIT: AI444613.1, EVALUE 8.50e-01
OTHER INFORMATION: SWISSPROT HIT: Q01851, EVALUE 9.20e+00
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                                                                                  APPLICATION NUMBER: US 09/774,203
                                                                                                          FILING DATE:
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
                                                                                                                      APPLICATION NUMBER: US 09/608,408
                                                                                                                                                           APPLICATION NUMBER: US 60/234,687
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FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
ADDITCATION NUMBER: PCT/US01/00663
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FILING DATE: 2001-01-30
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                    Annomax Sequence Listing Engine vers. 1.1
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Hanzel, David K.
Chen, Wensheng
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Pred. No. 2.8e+03;
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                                                            PRIOR APPLICATION NUMBER: US 60/234,687
                                                                                           PRIOR FILING DATE: 2001-01-30
                                                                                                          PRIOR APPLICATION NUMBER: PCT/US01/00670
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                                                PRIOR FILING DATE:
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PRIOR APPLICATION NUMBER: PCT/US01/00665
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
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TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Aeomica-X-1
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OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.1
OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.1
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 0.77
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 0.88
OTHER INFORMATION: EST. HUMAN HIT: BF475330.1, EVALUE 9.50e-01
OTHER INFORMATION: NT HIT: Z99119.1, EVALUE 1.60e-01
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            APPLICATION NUMBER: US 09/608,408
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                                                                                                                                                                                                                                                                                    FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: PCT/US01/00668
                                                                                                                                                                                        FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
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Chen, Wensheng
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2000-06-30
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OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.1
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.2
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.2
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.1
OTHER INFORMATION: EXPRESSED IN BADULT LIVER, SIGNAL = 1.1
OTHER INFORMATION: EXPRESSED IN BADULT LIVER, SIGNAL = 1.1
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.1
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.1
OTHER INFORMATION: MITHIT: AF303225.1, EVALUE 3.10e+00
US-09-864-761-28328
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US-09-864-761-25222
                            PRIOR APPLICATION NUMBER: PCT/USO1/0067
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00664
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR APPLICATION NUMBER: PCT/USO1/00665
PRIOR APPLICATION NUMBER: PCT/USO1/00665
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR APPLICATION NUMBER: PCT/USO1/00663
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PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
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APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
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PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1202 ATCCTAGGG 1210
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1142 AAGGAGAATTAAAGGAGTTGGAAAGTAGTACGGATGAGAGTGAAGAAGAACAAATCTCTG 1201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28 AGGCTAAGG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88 ACGGAGAAGTAATGGAATATAAGACTGGACAGCAAGGACGTGATTATGAAGAAATGTGTC 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
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FILING DATE: 2001-01-30
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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, OTHER INFORMATION: sandwich assay target and signaling probe US-09-847-113-6 \,
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                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/201,026
PRIOR FILING DATE: 2000-05-01
PRIOR PPLICATION NUMBER: 09/626,096
PRIOR APPLICATION NUMBER: 09/626,096
PRIOR FILING DATE: 2000-07-26
NUMBER OF SEQ ID NOS: 9
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 25222
LENGTH: 99
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Best Local
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PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR PRIOR DATE: 2000-09-21
PRIOR PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
PRIOR FILING DATE: 2000-06-30
                                                                                                                                                      Matches
                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Yu, Changjun TITLE OF INVENTION: Target Analyte Detection Using Asymmetrical Self-Assembled Mon FILE REFERENCE: A-69306-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/847,113
CURRENT FILING DATE: 2001-05-01
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OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 2.8

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 2.5

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 6.3

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 7.8

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 2.3

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 2.3

OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 6.7

OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 6.7

OTHER INFORMATION: EXT. HUMAN HIT: BE304683.1, EVALUE 4.00e-28

OTHER INFORMATION: SWISSPROT HIT: Q89940, EVALUE 5.50e+00

OTHER INFORMATION: NT HIT: AL163210.2, EVALUE 5.00e-30
                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 86
3931 GAAGAGG 3937
                                                                                                 3871 GTGGAGCTCCATAGAGAAAGAAATGTACATCTTCATGGAGTACTGCGATGAGGGGACTTTA 3930
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            924 GAAGGACTATGAAAAATATGCTGCAAAAGACTTCCAGGACAGGGTGCAGGCAC 976
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match 0.5%;
Local Similarity 66.0%;
                                               5 GTGGGGGGACATCAAGCAGCCATGCAAATGTTAAAAGAGACCATCAATGAGGAAGCTGCA 64
                                                                                                                                                                              Local
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                                                                                                                                                                                Similarity
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                                                                                                                                                         Conservative
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Pred. No. 3.2e+03;
0; Mismatches 18;
                                                                                                                                                                              Score 23.8; DB 10;
Pred. No. 3.7e+03;
                                                                                                                                                         Mismatches
                                                                                                                                                            27;
                                                                                                                                                                                                        Length 86;
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US-09-864
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                                                                                                                      OTHER
OTHER
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SOFTWARE: Annomax Sequence
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PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
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APPLICANT: Hanzel, David R.
APPLICANT: Chen, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Accomica-X-1
                                                                                                                                                                                                              OTHER
OTHER
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OTHER INFORMATION:
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CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
                                                           OTHER
                                                                                        OTHER
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                                                                                                                                                                                    OTHER
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                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 09/608,408 FILING DATE: 2000-06-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21
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FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00668
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: PCT/US01/00670
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APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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                      INFORMATION:
                                                                                                                                                INFORMATION:
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                                                                                        INFORMATION:
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                                                       INFORMATION:
                                                                                                                      INFORMATION:
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         ON: MAP TO ACO04613.1

NI EXPRESSED IN LUNG, SIGNAL = 2.2

NI EXPRESSED IN REAIN, SIGNAL = 1.7

DN: EXPRESSED IN ADULT LIVER, SIGNAL = 2

NI EXPRESSED IN PLACENTA, SIGNAL = 1.8

NI EXPRESSED IN HELA, SIGNAL = 2

NI EXPRESSED IN HELA, SIGNAL = 2

NI EXPRESSED IN HELA, SIGNAL = 1.7

NI EXPRESSED IN HONE MARROW, SIGNAL = 1.8

NI EXPRESSED IN FETAL LIVER, SIGNAL = 1.8
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NUMBER OF SEQ ID NOS: 2001-11-16;
SOFTWARE: Corixa Invention Disclosure Database;
SEQ ID NO 1313
LENGTH: 79
TYPE: DNA
ORGANISM: Homo sapiens
US-09-998-598-1313
       Db
                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-998-598-1313/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 12
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                                                                      Matches
                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1313, Application US/09998598 Patent No. US20020150922A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 4593
SEQ ID NO 453
                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 453, Application US/09969373 Patent No. US20020133852A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                           APPLICANT: Meagher, Madelein Joy
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.561
                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 37; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Effertz, Roger J.
APPLICANT: Hauge, Barian M.
TITLE OF INVENTION: Soybean SSRs and Methods of Genotyping
FILE REFERENCE: 38-10(52679)A
CURRENT APPLICATION NUMBER: US/09/969,373
CURRENT FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: US 09/754,853
PRIOR FILING DATE: 2001-01-05
PRIOR FILING DATE: 2001-01-05
                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Stolk, John A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 09/855,768 PRIOR FILING DATE: 2001-05-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 2001-01-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 09/760,427
                 3629 CCTATGATAATGTTATGCACGTTGGCTTGAGGAGGTGACCTTCAAATGGCAAAGAGGAA 3688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1106 AAGGTAATGAGCCGGAGTATGAGGGTGATGACACAGAAGGAGGAGATTAAAGGAGTTGGAA 1164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  188 AAACAGTGGAGAATGTGGAAGAATACAGCTATAAGCAGGAGAA 230
   8
                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29 AAGAAGTGGTTTATGAGGAAGAAGAAGCAAGAAGAAGAAGAA 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31,
                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                         Xu, Jiangchun
Chenault, Ruth A.
                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                 0.5%;
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                                                                                 Score 23.6; DB Pred. No. 4e+03;
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Pred. No. 4.1e+03;
0; Mismatches 12;
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Pred. No. 3.8e+03;
                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                 DB 10;
                                                                   26;
                                                     Indels
                                                                                                 Length 79;
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3689 ACAAAA 3694

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                                                              US-09-728-445-336
Sequence 336, Application US/09728445
Patent No. US20020102543A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Rosen, Craig A.
APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Human Genes, Sequences, and Expression Products 16.2
FILE REFERENCE: PO-16.2C1
CURRENT APPLICATION NUMBER: US/09/783,590
CURRENT FILING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 08/420,856
PRIOR APPLICATION NUMBER: 08/346,731
PRIOR FILING DATE: 1994-11-21
NUMBER OF SEQ ID NOS: 12485
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 9797
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Dillon, Patrick J.
APPLICANT: Haseltine, William A.
APPLICANT: Li, Haodong
APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
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  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc feature
LOCATION: (31)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (32)
OTHER INFORMATION: n equals a,t,g, or c
                                         APPLICANT: Friedrich, Glenn
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                    APPLICANT:
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OTHER INFORMATION: n equals a,t,g, or c
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LOCATION: (81)
OTHER INFORMATION: n equals a,t,9, or c
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                                                                                                                                                                                                                                                                                                                                                     498 AAAGCTTACCTCAGTCTCAAAGAAAAAAAGACAGGGAGCAAAGAGAGAAAATACGTC 557
                                                                                                                                                                                                                                                                       558 TGGTTTCTGGCTTA 571
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                                                                                                                                                                                                                                                                                                              4 AGAGTGAACCTCAGTAAAAAAAAAAAAANNAAAAGGAAAGAAAAAAAGATGNTAATTATATC 63
Sands, Arthur T.
                    Zambrowicz
                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                              33; Indels
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; TYPE: DNA; ORGANISM: Mus musculus US-09-728-445-336
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                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No. US20020013958A1 700159608H2
NAME/KEY: unsure
LOCATION: 43, 50, 52-53, 61, 63, 67, 69
OTHER INFORMATION: a, t, c, g, or other
US-09-923-876-1834
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CURRENT APPLICATION NUMBER: US/09/728,445
CURRENT FILING DATE: 2000-11-30
PRIOR APPLICATION NUMBER: US 60/168,358
PRIOR FILING DATE: 1999-12-01
NUMBER OF SEC ID NOS: 891
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SEQ ID NO 1834
                                                                                                                                                                                        Matches
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CURRENT FILING DATE: 2001-08-06
PRIOR APPLICATION NUMBER: 09/298,329
PRIOR FILING DATE: 1999-04-21
PRIOR APPLICATION NUMBER: 60/085,331
PRIOR PILING DATE: 1998-05-05
PRIOR PILING DATE: 1998-05-05
NUMBER OF SEQ ID NOS: 6332
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APPLICANT: Kamigaki, Laura Y. (Ito)
APPLICANT: Sherman, Bradley K.
APPLICANT: Sherman, Bradley K.
TITLE OF INVENTION: POLYMUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN SEEDLING
FILE REFERENCE: PL-0012-1 CON
                                                                                                                                                                                                                                  Query Match
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                                                                                        486 ACGACTTTTGCTAAAGCTTACCTCAGTCTCAAAGAAAAAGACAGGGAGCAAAGAGGGACA 545
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                                              546 AGAAA 550
66 ANANA 70
                                                                                                                                                                                                              Local
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US-09-783-590-9597/c

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US-09-864-761-20158
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: GB 24263.6
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SEQ ID NO 9597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Hanzel, David | APPLICANT: Chen, Wensheng
                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/180,312
                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/864,761 CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: Aeomica-X-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
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PRIOR FILING DATE: 1994-11-21
NUMBER OF SEQ ID NOS: 12485
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/783,590 CURRENT FILING DATE: 2000-02-15 PRIOR APPLICATION NUMBER: 08/420,856 PRIOR FILING DATE: 1995-04-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: n equals a,t,g, or c
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ORGANISM: Homo sapiens
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1967 AAAATCTGTTAGAAGAAGAATGGAATTTCACCAAAGAAATAACTCATTACATA 2019
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 64.2%;
                                                  APPLICATION NUMBER: PCT/US01/00665
                                                                    APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
                                                                                                          APPLICATION NUMBER: PCT/US01/00664 FILING DATE: 2001-01-30
                                                                                                                                                APPLICATION NUMBER: PCT/US01/00667 FILING DATE: 2001-01-30
                                                                                                                                                                                                         APPLICATION NUMBER: PCT/US01/00666
                                                                                                                                                                                                                         APPLICATION NUMBER: US 60/236,359 FILING DATE: 2000-09-27
                                                                                                                                                                                                                                                                   FILING DATE: 2000-10-04
            APPLICATION NUMBER: PCT/US01/00668
                                    FILING DATE:
                                                                                                                                                                                           FILING DATE:
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D. US20020110850A1
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Hanzel, David K.
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Ruben, Steven M.
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2001-01-30
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Pred. No. 5.7e+03;
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; ORGANISM: Murine US-09-933-797-749
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OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 5.1

OTHER INFORMATION: EXPRESSED IN BTA74, SIGNAL = 5.1

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 5.5

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 5.5

OTHER INFORMATION: EXPRESSED IN HELLO, SIGNAL = 5.2

OTHER INFORMATION: EXPRESSED IN HBL100, SIGNAL = 4.8

OTHER INFORMATION: EXPRESSED IN ADJUT LIVER, SIGNAL = 5.8

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 5.8

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 4.3

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 4.3

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 4.6

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 4.6

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 4.6

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OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 4.6

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 4.6

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 4.6

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 4.6

OTHER INFORMATION: MISSPROT HIT: 054828, EVALUE 7.00e+00

THER INFORMATION: MISSPROT HIT: D54828, EVALUE 7.70e-01
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Query Match
Best Local Similarity
                                                                                                                                                                                                                            SOFTWARE: Fas
SEQ ID NO 749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 749, Application US/09933797 Patent No. US20020155119A1
                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Robert A. Sikes et al.

TITLE OF INVENTION: Isolation and Use of Fetal Urogenital

TITLE OF INVENTION: Sinus Expressed Sequences

FILE REFERENCE: 9901-007-999

CURRENT APPLICATION NUMBER: US/09/933,797

CURRENT FILING DATE: 2001-08-22

PRIOR APPLICATION NUMBER: US/09/482,933

PRIOR APPLICATION NUMBER: US/09/482,933

PRIOR FILING DATE: 2000-01-14
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                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 1999-05/14
PRIOR APPLICATION NUMBER: 60/085,383
                                                                                                                                                                                                                                                                                            NUMBER OF SEQ
                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 1998-05-14
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                                                                                                                                                                  TYPE: DNA
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PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
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PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
                                                                                                                                                                                                  LENGTH: 94
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
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PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                     FastSEQ for Windows Version 3.0
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53.7%;
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Pred. No. 5.8e+03;
0; Mismatches 23;
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Pred. No. 5.9e+03;

Score

23.2;

DB 9;

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Patent No. US20020048763A1
GENERAL INFORMATION:
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CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 2000-08-03
                                                                                                                                                                                                                                                 SOFTWARE: Annomax Sequence Listing Engine vers. 1.1 SEQ ID NO 27858
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Hanzel, David APPLICANT: Chen, Wensheng
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                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
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OTHER INFORMATION: MAP TO AC005386.1
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 0.68
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.79
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.86
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 0.76
                                                                                                                                             FEATURE:
                                                                                                                                                                ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OR APPLICATION NUMBER: PCT/USO1/00667
OR FILING DATE: 2001-01-30
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/USO1/00664
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/USO1/00665
OR APPLICATION NUMBER: PCT/USO1/00665
OR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-90 APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21
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US-09-864-761-32598/c
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                               PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 32598
LENGTH: 96
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CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 2000-08-03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00664

PRIOR FILING DATE: 2001-01-20
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PRIOR APPLICATION NUMBER: PCT/US01/00667
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ORGANISM: Homo sapiens
                      TYPE: DNA
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FILING DATE: 2000-06-
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FEATURE:

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US-09-864-761-25187; Sequence 25187, Application US/09864761
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SEQ ID NO 25187
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                                                                                                                                                                         NUMBER OF SEQ ID NOS: 49117
                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 09/774,203 PRIOR FILING DATE: 2001-01-29
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OTHER INFORMATION: MAP TO ACO11449.3
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 2.5
                                                     ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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                                                                                               TYPE: DNA
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PRIOR FILING DATE: 2001-01-30
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                                                                                                              LENGTH: 97
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FILING DATE: 2000-06-30
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INFORMATION: EXPRESSED IN BONE
INFORMATION: SWISSPROT HIT: P1
                                                                                                                                              Annomax Sequence Listing Engine vers. 1.1
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SWISSPROT HIT: P17588, EVALUE 2.50e+00
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RESULT 22
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                                                                               APPLICATION NUMBER: FILING DATE: 2001-0
                                                         APPLICATION NUMBER: US 60/234,687
                                                                                                       APPLICATION NUMBER: PCT/US01/00661 FILING DATE: 2001-01-30
              APPLICATION NUMBER: US 09/608,408 FILING DATE: 2000-06-30
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                                                                                                                                       APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
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APPLICATION NUMBER:
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Chen, Wensheng
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EXPRESSED IN LUNG, SIGNAL = 3.3
EST_HUMAN HIT: BE393813.1, EVALUE 9.00e-48
NT HIT: M28624.1, EVALUE 5.20e-02
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09/774,203
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IN BRAIN, SIGNAL = 3.1
IN PLACENTA, SIGNAL = 3.1
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IN HELA, SIGNAL = 3.3
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Pred. No. 6
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                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR PRIOR DATE: 2000-10-04
PRIOR PPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 700/666
PRIOR PILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US 700/666
PRIOR PRILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: DCT/US01/00667
PRIOR APPLICATION NUMBER: DCT/US01/00667
PRIOR FILING DATE: 2001-01-30
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Best Local (
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APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Acomica-X-1
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CURRENT FILING DATE: 2001-05-23
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PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR PRIOR DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
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INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.9
INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 2.8
INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 3.5
INFORMATION: EXPRESSED IN LUNG, SIGNAL = 3.5
INFORMATION: EXPRESSED IN HEART, SIGNAL = 2.5
INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 3.8
INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 3.8
INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 3.8
INFORMATION: MT HIT: 9111434115, EVALUE 9.00e-25
INFORMATION: SWISSPROT HIT: P044424, EVALUE 9.00e-07
INFORMATION: SWISSPROT HIT: P044424, EVALUE 9.00e-07
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David K.
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pred. No. 6.2e+03;
pred. No. 6.3e+03;
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FEATURE:

OTHER INFORMATION: AD TO AC009401.1

OTHER INFORMATION: EXPRESSED IN BH474, SIGNAL = 1.3

OTHER INFORMATION: EXPRESSED IN BH474, SIGNAL = 1.3

OTHER INFORMATION: EXT_HUMAN HIT: AW237754.1, EVALUE 3.00e-03

OTHER INFORMATION: SWISSPROT HIT: 015390, EVALUE 5.40e+00

OTHER INFORMATION: NT HIT: AF092030.2, EVALUE 4.30e-02

US-09-864-761-33084
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PRIOR PELLON NUMBER: US 60/207,456
PRIOR PELLON NUMBER: US 60/207,456
PRIOR PELLON NUMBER: US 09/632,366
PRIOR PELLON NUMBER: US 2006-08-03
PRIOR PELLON NUMBER: US 2006-08-03
PRIOR PELLON NUMBER: US 60/236,359
PRIOR PELLON NUMBER: US 60/236,359
PRIOR PELLON DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
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NUMBER OF SEQ ID NOS: 49117

SOFTWARE: Annomax Sequence Listing Engine vers.
SEQ ID NO 33084

LENGTH: 84
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PRIOR EILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 99/608,408
PRIOR PILING DATE: 2000-06-30
PRIOR PILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 09/774,203
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION UNMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY FILE REFERENCE: Acomica-x-1 File Reference: Acomica-y-1-761 CURRENT APPLICATION NUMBER: US/09/864,761 CURRENT FILING DATE: 2001-05-23 CURRENT FILING DATE: 2001-05-23
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APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K
APPLICANT: Chen, Wensheng
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                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: ZUUL-VI-VI
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR APPLICATION 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
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Pred. No.
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US-09-864-761-27695/c
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PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR APPLICATION UMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR APPLICATION NUMBER: US 09/608,408
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Patent No. US20020048763A1
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PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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NUMBER OF SEQ ID NOS: 49117
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OTHER INFORMATION: NT HIT: L17000.1, EVALUE 1.00e-33
OTHER INFORMATION: SWISSPROT HIT: Q16566, EVALUE 1.00e-08
OTHER INFORMATION: EST_HUMAN HIT: AL134969.1, EVALUE 1.00e-33
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PRIOR FILING DATE: 2001-01-30
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                                               APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
                                                                                                                               FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
                                                                                                                                                                                APPLICATION NUMBER: PCT/US01/00664 FILING DATE: 2001-01-30
FILING DATE: 2001-01-30
                FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 60/236,359
FILING DATE: 2000-09-27
APPLICATION NUMBER: PCT/US01/00666
                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                      APPLICATION NUMBER: PCT/US01/00667
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                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: GB 24263.6
FILING DATE: 2000-10-04
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 09/632,366 FILING DATE: 2000-08-03
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Chen, Wensheng
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Pred. No.
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US-09-864-761-31482/c
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                                                                             PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 09/632,366 PRIOR FILING DATE: 2000-08-03 PRIOR APPLICATION NUMBER: GB 24263.6
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PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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                                            PRIOR FILING DATE:
                                                                                                                                                                             PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/864,761 CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Penn, Sharron G. APPLICANT: Rank, David R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 49117
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
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OTHER INFORMATION:
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                  FILING DATE: 2001-01-30
PTITING DATE: 2001-01-30
PTITING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00664 FILING DATE: 2001-01-30
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Hanzel, David K.
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DN: EXPRESSED IN HUNG, SIGNAL = 0.98

DN: EXPRESSED IN BARAN, SIGNAL = 1.1

DN: EXPRESSED IN BONE MARROW, SIGNAL = 0.93

DN: EXPRESSED IN FUTAL LIVER, SIGNAL = 0.93

DN: EXT_HUMAN HIT: AI797128.1, EVALUE 2.90e-01

DN: NT HIT: X95199.1, EVALUE 3.10e+00
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Pred. No. 6.8e+03;
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OTHER INFORMATION: MAP TO AC007567.1

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.7

OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 2.3

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.4

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.4

OTHER INFORMATION: EXPRESSED IN LONG, SIGNAL = 1.2

OTHER INFORMATION: NT HIT: AL163303.2, EVALUE 1.00e-05

OTHER INFORMATION: EST_HUMAN HIT: BE221737.1, EVALUE 1.00e-03

OTHER INFORMATION: SWISSPROT HIT: P13186, EVALUE 1.10e+00

US-09-864-761-31482
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US-09-864-761-32824/c
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                                                                                                           CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR PRIOR DATE: 2000-08-03
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
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Best Local :
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PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR APPLICATION NUMBER: US 09/774,203
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION UMMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR APPLICATION NUMBER: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION UNMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY FILE REFERENCE: Aeomica-X-1
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PRIOR APPLICATION NUMBER: PCT/US01/00666 PRIOR FILING DATE: 2001-01-30
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                                                          APPLICATION NUMBER: US
FILING DATE: 2000-09-27
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Chen, Wensheng
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                                                                                             US 60/236,359
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; OTHER INFORMATION: MAP TO AL030997.1
; OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 0.93
; OTHER INFORMATION: NT HIT: M96655.1, EVALUE 1.80e-01
US-09-864-761-32824
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SEQ ID NO 32824
LEBUGTH: 89
TYPE: DNA
                                                                                                                                                                           NUMBER OF SEQ ID NOS: 6332
SOFTWARE: PERL PROGRAM
SEQ ID NO 1300
LENGTH: 98
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Best Local :
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION UNMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
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APPLICANT: Kamigaki, Laura Y. (Ito)
APPLICANT: Sherman, Bradley K.
TITLE OF INVENTION: POLYMUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN SEEDLING
FILE REPERROE: PL-0012-1 CON
CURRENT APPLICATION NUMBER: 09/293,876
CURRENT FILING DATE: 109/2983.329
PRIOR APPLICATION NUMBER: 09/2983.329
PRIOR PILICATION NUMBER: 60/2985.331
PRIOR PILICATION NUMBER: 60/2985.331
PRIOR PILICATION NUMBER: 60/2985.331
PRIOR FILING DATE: 1998-05-05
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PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
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PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR BETTING PRIOR PR
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION UNMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00670
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                                                                              TYPE: DNA
ORGANISM: Zea mays
NAME/KEY: misc_feature
                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2041 GGGAAGCTTTTCTG 2054
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26 CGTAGACTTATCTG 13
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                                                                                                                                                                                              Sequence 7267, Application US/09878574 Patent No. US20020110548A1
     TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with TITLE OF INVENTION: Plants FILE OF INVENTION: Plants FILE REFERENCE: 38-21(15401)B CURRENT APPLICATION NUMBER: US/09/878,574 CURRENT FILING DATE: 2001-12-21 PRIOR APPLICATION NUMBER: 09/333,535 PRIOR FILING DATE: 199-06-14
                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 637
LENGTH: 64
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                 -09-878-574-7267/c
NUMBER OF SEQ ID NOS: 15775
                                                                                                                                    APPLICANT:
                                                                                                                                                  APPLICANT:
                                                                                                                                                              APPLICANT: Byrum, Joseph R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 637,
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APPLICANT: Xu, Jiangchun
APPLICANT: Secrist, Heat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/919,580 CURRENT FILING DATE: 2001-07-30
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TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: n = A, T, C \text{ or } G
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                                                                                                                             La Rosa, Thomas J. Thompson, Michael D.
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Pred. No. 6.6e+03;
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Pred. No. 7.9e+03;
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US-09-864-761-24215
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                      NUMBER OF SEQ ID NOS: 49117
                                     PRIOR FILING DATE:
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Chen, Wensheng
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                                     2001-01-29
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; ORGANISM: Glycine max
; OTHER INFORMATION: Clone ID: 701099674H1
US-09-878-574-7267
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1 SEQ ID NO 24215
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                                                                                                  PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 09/774,203
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PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
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PRIOR APPLICATION NUMBER: US 60/236,359
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4593 GCACAGGTGACAAGCGTCACTTCTC 4617
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                                                                                                                                                                                                                                                       APPLICATION NUMBER: PCT/US01/00670
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: PCT/US01/00665
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                                                                                                                                                     APPLICATION NUMBER: US 09/608,408
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Pred. No. 8.9e+03;
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RESULT 32
US-09-864-761-25667
; Sequence 25667, Application US/09864761
: Datent No. US20020048763A1
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                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 2000-05-26

PRIOR APPLICATION UNMBER: US 09/632,366

PRIOR FILING DATE: 2000-08-03

PRIOR PILING DATE: 2000-08-03

PRIOR PILING DATE: 2000-08-03

PRIOR FILING DATE: 2000-08-03

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR FILING DATE: 2000-09-27

PRIOR PPLICATION NUMBER: PCT/US01/00666

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00664

PRIOR PILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00669

PRIOR APPLICATION NUMBER: PCT/US01/00665

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30
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APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
APPLICANT: Chen, Wensheng
APPLICANT: Chen, Wensheng
APPLICANT: Chen, Wensheng
APPLICANTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILLE REFERENCE: Acomica-x-1
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR FILING DATE: 2000-02-04
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Local Similarity 58.0%;
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INFORMATION: EXPRESSED IN PLACEWTA, SIGNAL = 1.3
INFORMATION: EXPRESSED IN FETAL LIVEE, SIGNAL = 1.5
INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 0.87
INFORMATION: EXPRESSED IN ADULT LIVEE, SIGNAL = 1.2
INFORMATION: EXPRESSED IN HEART, SIGNAL = 1.2
INFORMATION: EXPRESSED IN LONG, SIGNAL = 1.2
INFORMATION: EXPRESSED IN ENERGY, SIGNAL = 1.9
INFORMATION: EXPRESSED IN ENERGY, SIGNAL = 1.9
INFORMATION: EXPRESSED IN ENERGY, SIGNAL = 1.9
INFORMATION: EXPRESSED IN ENERGY SIGNAL = 1.9
INFORMATION: EXPRESSED IN ENERGY SIGNAL = 1.9
INFORMATION: NT HIT: M36383.1, EVALUE 3.20e+00
APPLICATION NUMBER: US 60/234,687
                                                                                         APPLICATION NUMBER: PCT/US01/00662 ETILING DATE: 2001-01-30 APPLICATION NUMBER: PCT/US01/00661 FILING DATE: 2001-01-30
                                                                                                                                                                                                                         APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
                                     APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30
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Pred. No. 9
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FEATURE:

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US-09-878-574-14838
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PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 25667
LEBGTH: 100
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APPLICANT: Byrum, Joseph R.

APPLICANT: La Rosa, Thomas J.

APPLICANT: Thompson, Michael D.

APPLICANT: Thompson, Michael D.

APPLICANT: Thompson, Michael D.

APPLICANT: Numerion: Acid Molecules and Other Molecules Associated with
TITLE OF INVENTION: Plants

TITLE OF INVENTION: Plants

FILE REFERENCE: 38-21(15401)B

CURRENT APPLICATION NUMBER: U5/09/878,574

CURRENT APPLICATION NUMBER: 09/333,535

PRIOR APPLICATION NUMBER: 09/333,535
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LENGTH: 83
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Glycine max
OTHER INFORMATION: Clone ID: 701069318H1
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                                                                               3237 GAAC 3240
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66 GATC 69
                                                                                                                                                                    6 TGGTCGTGTCGCAGCAAAACCAAGCCATCCACTAACTTGTTCTGTCCAAGACATGGCTCT 65
                                                                                                                                                                                                                                                                                                                                                                                 Match 0.4%;
Local Similarity 59.4%;
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Pred. No. 9.1e+03;
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US-09-864-761-19403
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                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Annomax Sequence Listing Engine vers. SEQ ID NO 19403
                                                                                                    OTHER
OTHER
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OTHER INFORMATION:
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PRIOR APPLICATION NUMBER: PCT/US01/00667
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PRIOR APPLICATION N
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RINFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.4
RINFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.4
RINFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.4
RINFORMATION: EXPRESSED IN HELA, SIGNAL = 1.5
RINFORMATION: EXPRESSED IN HELA, SIGNAL = 1.4
RINFORMATION: EXPRESSED IN HELA, SIGNAL = 1.4
RINFORMATION: SWISSPROT HIT: Q61151, EVALUE 8.00e-12
RINFORMATION: EST-HUMAN HIT: BE929768.1, EVALUE 3.00e-761-19403
                                                                                                                                                                           INFORMATION:
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APPLICATION NUMBER: US 09/608,408
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Hanzel, David K.
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                                                                                                                                                                                                                                         IN BONE MARROW, SIGNAL = 1.8
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                                                                                                                                                                                                       BT474, SIGNAL = 1.4
FETAL LIVER, SIGNAL = 1.1
                                                                                                                                                                                       HEART, SIGNAL = 1.7
                                EVALUE 3.00e-44
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OTHER INFORMATION: MAP TO AC005666.1
OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.1
OTHER INFORMATION: EXPRESSED IN LING, SIGNAL = 1.1
OTHER INFORMATION: EXPRESSED IN ETTAL LIVER, SIGNAL = 0
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 5.6
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 0
OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1
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Best Local Similarity 62.5
Matches 35; Conservative
                                                                                                                                                                                                                               SOFTWARE: Annomax
                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 49117
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PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
                                                                                                                                                    ORGANISM: Homo sapiens
                                                                                                                                                                              TYPE: DNA
                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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                                                                                                                                                                                                                                                                                FILING DATE: 2000-06-30
APPLICATION NUMBER: US 09/774,203
                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21
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APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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Hanzel, David K.
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Pred. No. 1e+04;
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US-09-864-761-27595
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US-09-864-761-24218/c
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PRIOR PELICATE ON UNMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PELICATION NUMBER: US 09/632,366
PRIOR PELICATION NUMBER: US 09/632,366
PRIOR PELICATION NUMBER: US 09/632,366
PRIOR PELICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: DCT/US01/00666
PRIOR APPLICATION NUMBER: DCT/US01/00666
PRIOR APPLICATION NUMBER: DCT/US01/00667
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Patent No. US20020048763A1
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                                                                                                                                                        SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 24218
LENGTH: 75
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APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
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CURRENT FILING DATE: 2001-05-23
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OTHER INFORMATION: NT HIT: U88870.1, EVALUE 1.00e-49
OTHER INFORMATION: EST_HUMAN HIT: AW964640.1, EVALUE 1.00e-49
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PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 09/608,408 PRIOR FILING DATE: 2000-06-30
OTHER INFORMATION: MAP TO AC013746.4
OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 3.8
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                                                                FEATURE:
                                                                                     ORGANISM: Homo sapiens
                                                                                                                        TYPE: DNA
                                                                                                                                                     ENGTH:
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FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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NT HIT: U88870.1, EVALUE 1.00e-49
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Pred. No. 1
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US-09-969-373-613/c
Sequence 613, Application US/09969373
Patent No. US20020133852A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-864-761-22604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 38
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PRIOR FILING DATE: 2001-05-15
NUMBER OF SEQ ID NOS: 4593
SEQ ID NO 613
LENGTH: 80
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Best Local Similarity
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APPLICANT: Hauge, Brian M.

APPLICANT: Hauge, Brian M.

APPLICANT: Hauge, Brian M.

APPLICANT: Hauge, Brian M.

APPLICANT: ABUREN: Soybean SSRs and Methods of Genotyping FILE REFERENCE: 38-10(52679)A

CURRENT APPLICATION NUMBER: US/09/969,373

CURRENT FILING DATE: 2001-10-02

CURRENT FILING DATE: 2001-10-02
                               APPLICANT: Renn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
APPLICANT: Chen, Wensheng
APPLICANT: Chen, Gene Expression Analysis by Microarray
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
TILLE REFERENCE: Accomica-x-1
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 22604, Application US/09864761
Patent No. US20020048763A1
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PRIOR FILING DATE: 2001-01-05
PRIOR APPLICATION NUMBER: US 09/760,427
PRIOR FILING DATE: 2001-01-13
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PRIOR FILING DATE: 2000-05-26
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PRIOR APPLICATION NUMBER: US 09/632,366 PRIOR FILING DATE: 2000-08-03

APPLICATION NUMBER: GB 24263.6

FILING DATE:

FILING DATE:

APPLICATION NUMBER: US 60/236,359

2000-10-04 NUMBER: US 1 2000-09-27

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US-09-878-574-5668

: Sequence 5668, Application US/09878574

: Patent No. US20020110548A1
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APPLICANT: ia Rosa, Thomas J.
APPLICANT: Thompson, Michael D.
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(15401)B
CURRENT APPLICATION NUMBER: US/09/878,574
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                                                                                                                                           APPLICANT: Byrum, Joseph R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.7
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.9
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.9
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.6
OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.8
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 1.8
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.7
OTHER INFORMATION: EXPRESSED IN BEANT, SIGNAL = 1.7
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.7
OTHER INFORMATION: NT HIT: G1103653, EVALUE 6.70e-01
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PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
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ORGANISM: Homo sapiens
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APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
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Pred. No. 1e+04;
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Search completed: November 11, 2002, 12:43:10 Job time: 136 secs
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EARLIER FILING DATE: 1995-01-17
EARLIER APPLICATION NUMBER: US 08/353,537
EARLIER APPLICATION NUMBER: US 08/353,537
EARLIER APPLICATION NUMBER: POT/US94/05908
EARLIER FILING DATE: 1993-05-25
EARLIER APPLICATION NUMBER: US 08/110,286
EARLIER FILING DATE: 1993-08-23
EARLIER APPLICATION NUMBER: US 08/079,320
EARLIER FILING DATE: 1993-06-18
NUMBER OF SEQ ID NOS: 15
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                                                                                                                                                                                           Query Match
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LENGTH: 86
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APPLICANT: Vale Jr., Wylie W.
APPLICANT: Donaldson, Cynthia J.
APPLICANT: Sawchenko, Paul
                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/191,724
CURRENT FILING DATE: 1998-11-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Cloning and Recombinant Production of TITLE OF INVENTION: CRF Receptor(s)
FILE REFERENCE: Salk1748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Perrin, Marilyn H. APPLICANT: Chen, Ruoping
                                                                                                                                                                                                                                                                        NAME/KEY: CDS
LOCATION: (1)
                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 15775
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                                                                                                                                                                                                                                                      OTHER INFORMATION:
                                                                                           4620 GCTGCTCTGTTTGTCTGATGTGGCAAAAGGCCCTCTGGAGGGCTGGTGGCCACGAGGT 4678
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ORGANISM: Glycine max
OTHER INFORMATION: Clone ID: 701097233H1
                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 87
                                                                                                                                                                       y Match 0.4%;
Local Similarity 61.0%;
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                                                                                                                                                                  Score 22.2; DB 10; Length 87; Pred. No. 1.1e+04;
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IMAGE:1361691 5', mRNA sequence.
                        Unpublished (1996)
Washington University School of MedicineP
                               The WashU-HHMI Mouse EST Project
                                         Waterston, R.
                                                                                          Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                         Mus musculus
                                                                                                                           EST
                                                                                                                                  AA990067.1 GI:3175431
                                                                                                                                          AA990067
                                                                                                                  house mouse.
                                                                                 (bases 1 to 95)
                                                                                                                                                                                                                                                ; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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D AV672913
7 AZ346768
BQ290138
AA918358
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AI312199
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BQ526570
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BF533538
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AZ346768 1M0082H10
BQ290138 hd74c10.g
AA918358 on42h01.s
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BE196765 SAC181 Mo
AZ920583 1006020E1
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AI312199 ta90a04.x
AI719156 at06d02.x
                                                                                                                                                                                                                                                                                                                            BQ526570 NISC_no17
BH866027 SALK_1003
AI309695 tb56al0.x
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AA761267 nz25c08.s
AA958438 ua12c12.r
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AI247119 gx52f10.x
BE636445 SWOVL2CA
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AZ920585 1006020E1
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AZ482078 1M0307B04
AZ784737 2M0027L14
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T61675 yb87g06.s1
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BE933241 RC1-HT088
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                                                                                                                                             ORGANISM
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Best Local
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                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes; Cyprinidae; Danio.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5α
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                                                                                                                 part of the Daniokey Pilot BAC Library created by R. Plasterk and
                                                                                                                                                                                     Submitted (06-JUN-2002) The Sanger Institute,
                                                                                                                                                                                                                                                                                                             Danio rerio
                                                                                                                                                                                                                                                                                                                                               AL743114.1
GSS.
                                                                                                                                                                                                          Direct Submission
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                                                                                                                                                                                                                                                                                                                                 zebrafish
                                                                                                                                                                                                                                                                                                                                                                                 AL743114
                                                                                                                                                                                                                                                                                                                                                                                        Danio rerio genomic clone DKey-12A12, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                   DRIZA12S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                  sequence was generated from the SP6 end of BAC 12A12. 12A12 is
                                                                                                  Keygene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               details: http://www.sanger.ac.uk/Projects/D_rerio/
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                /organism="Danio rerio"
/db_xref="taxon:7955"
/clone="DKey-12A12"
                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="IMAGE:1361691"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /tissue_type="Thymus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="Soares_thymus_2NbMT"
/sex="male"
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70.7%;
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Pred. No. 14;
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                                                                                                                                                                     CB10 1SA, UK. E-mail contact:
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DEFINITION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hiller, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost, S., Kucaba, T., Lacy, M., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B., White, Y., Wylie, T., Waterston, R. and Wilson, R.

Washu Merck EST Project 1997
Contact: Wilson RC (1997)
Contact: Wilson RC Walson, Wilson RC (1997)
Contact: Wilson RC (1997)
Conta
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 70)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: estewatson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Possible reversed clone: similarity on wrong strand
Seq primer; -28mil revz ET from Amersham
High quality sequence stop: 1.
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     19 a
/lab_host="BulbB"
/lab_host="DelibB"
/lab_host="DelibB"
/note="Organ: mixed (see below); Vector: pT7T3D-Pac
/note="Organ: mixed (see below); Vector: pT7T3D-Pac
/note="Organ: mixed (see below); Vector: p1smid DNA from tree
Site_2: Eco RI; Equal amounts of plasmid DNA from three
Site_2: Eco RI; Equal amounts of plasmid DNA from three
sommalized libraries (melanocyte 2NbHM, pregnant uterus
NbHPU, and fetal heart NbHHJ9W) were mixed, and ss circles
NbHPU, and fetal heart NbHHJ9W) were mixed, in this DNA
were made in vitro. Following HAP purification, this DNA
was used as tracer in a subtractive hybridization
was used as tracer in a subtractive hybridization
reaction. The driver was PCR-amplified cDNAs from pools of
5,000 clones made from the same 3 libraries. The pools
consisted of I.M.A.G.E. clones 260232-265223,
340488-345479, and 484488-489479."

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7 c 9 g 30 t
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/db_xref="taxon:9606"
/clone="IMACE:823794"
/clone="ib="Soares_NhHMPu_S1"
/tissue_type="Pooled human melanocyte, fetal heart, and pregnant uterus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 30; DB 17; Length 64; 
pred. No. 5.8e+03; 
0; Mismatches 20; Indels
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RESULT 4
AI767078
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AUTHORS
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Best Local Similarity 71.7%;
Matches 38; Conservative
                                       γ
                                                                                      В
                                                                                                                                                                                                                                                                                    BASE COUNT
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                                                                                                                                                                            Matches
                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4051 AATATCTTCCTTACCTCATCTGGATTAATCAAACTGGGAGATTTTGGATGTTC 4103
                                                                                                           3085 TTTGATTTTCTACAAGCAATTGAACCTGCCTTTATTTCAGCTTTTACCAGAAGAAGACTTC 3144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       source
                                         3145 TTGAGTT 3151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     62 TTGATTT 68
                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        71 bp mRNA linear EST 21-DEC-1999
wi92g07.x1 NCI_CGAP_Kid12 Homo sapiens cDNA clone IMAGE:2400828 3',
mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
cDNA sequencing by: Washington University Genome Sequencing Center
DNA sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 785 Std Error: 0.00
Seg primer: -40UP from Glbco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
Email: ggapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 71)
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         High quality sequence stop: 70.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1997)
                                                                                                                                                                            43;
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                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                            22 a
                                                                                                                                                                                                                                                                                                                    /note-Torgan: kidney; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: ECO RI; a modified polylinker; Site_1: Not I; Site_2: ECO RI; a modified polylinker; Site_1: Not I; Site_2: ECO RI; a plasmid DNA from the normalized library NOT_CGAP_Kid5 was plasmid DNA from the normalized in vitro. Following HAP purification, this DNA was used as tracer in a subtractive purification reaction. The driver was PCR-amplified CDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1323912-1325831, 1471368-1472903 and 1492104-1493255). Subtraction by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                          Fatima Bonaldo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone="IMAGE:2400828"
/clone_lib="NCI_CGAP_Kid12"
/tissue_type="2 pooled tumors (clear cell type)"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                 0.6%;
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Pred. No. 1.1e+04

    Mismatches

                                                                                                                                                                                                   Score 28.6; DB 9; Length 71; Pred. No. 1.5e+04;
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REFERENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      135 TTTGAAAGAAAAATGAATGCACCAAATCAGCCTCCACATAAAGACACTGGAAAAAACAGT 194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  75 TGNNAAAAAAAAATGAAACNAAAAAAAAAACATCCATTTTAAAACATTGTCACACAAGT 16
                                                                                                                                                                                                                                                                                                                                                                                                                      15 GGA 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,F., Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                         Arabidopsis thaliana
                                                                                                                                                                                                                                                            SALK_093795 Arabidopsis thaliana TDNA linear GSS 05-AUG-200 thaliana genomic clone SALK_093795, DNA sequence.
                                                                                                                                                                                                thale cress.
                                                                                                                                                                                                                                    BH863389.1 GI:22098952
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: tshini@genes.nig.ac.jp.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kitayama,A., Terasaka,C., Mochii,M., Ueno,N., Shin-i,T. and Kohara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fax: 81-559-81-6855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Tadasu Shin-i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Expressed genes in X. laevis embryo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BJ063432
BJ063432 NIBB Mochii normalized Xenopus tailbud library Xenopus taevis cDNA clone XL076e13 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  xenopodinae; xenopus.
1 (bases 1 to 83)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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                                                                                             (bases 1 to 65)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: pBSRN3; Site_1: Not1; Site_2: EcoR1; cDNAs were oligo-dT primed and directionally cloned. Staging according to Nieuwkoop and Faber. Library is subtracted and was constructed by N. Garrett and A.M. Zorn, Wellcome/CRC Institute). " 42 t 4 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /dev_stage="stage 25"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="NIBB Mochii normalized xenopus tailbud
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:8355"
/clone="XL076e13"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /tissue_type="whole_embryo"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Xenopus laevis"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.6%;
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Pred. No. 2.3e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 AAAATATATCATAGCAGGACAATTGAACCAGACCAAAAAAATAACAGCTTAAATA 55
                                                               Seq primer: -40UP from Gibco
High quality sequence stop: 76.
Location/Qualifiers
                                                                                                             www-bio.llni.gov/bbrp/lmage/lmage.html
Insert Length: 1558 Std Error: 0.00
Seg primer: -40UP from Gibco
                                                                                                                                                     CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1998)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                     Bonaldo, Ph.D.
                                                                                                                                                                                                                                                                                                               Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 83)
NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurological
Disorders and Stroke, Brain Tumor Genome Anatomy Project
                                                                                                                                                                                                                                                                                                                                                    Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                     cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima
                                                                                                                                                                                                                                                                                                                                                                                                                      (CGAP/BTGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9y89d09.x1 NCI_CGAP_Brn25 Homo sapiens cDNA clone IMAGE:2019185 3',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AI360096.1 GI:4111717
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mRNA sequence.
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Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Class: TDNA tagged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This is single pass sequence recovered from the left border of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           36 a
                  /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ecker@salk.edu
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 9 c 7 g 13 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone="SALK_093795"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:3702"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Arabidopsis thaliana"
/strain="Columbia 0"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                 Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 27.8; DB 17
Pred. No. 2.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         83 bp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 65;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EST 16-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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BASE COUNT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       535 CAAAGAGGACAAGAAAATA 553
                                                                                                                                                             source
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McCarter, J., Clifton, S., Chiapelli, B., Pape, D., Martin, J., Wylle, T., Dante, M., Marra, M., Hillier, L., Kucaba, T., Theising, B., Bowers, Y., Gibbons, M., Ritter, E., Bennett, J., Franklin, C., Tsagareishvili, R., Ronko, I., Kennedy, S., Maguire, L., Beck, C., Underwood, K., Steptoe, M., Allen, M., Person, B., Swaller, T., Harvey, N., Schurk, R., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R., Waterston, R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BI322285 94 bp mRNA linear EST 30-JUL-2001 kx19a08.y3 Parastrongyloides trichosuri FL pAMP1 v1 Chiapelli McCarter Parastrongyloides trichosuri cDNA 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Parastrongyloides trichosuri
Parastrongyloides trichosuri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The Washington Univ. Nematode EST Project, 1999
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Parastrongyloides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BI322285.1 GI:15001471
                                                                                                                                                                                                                                                                                                              Email: est@watson.wustl.edu
The library was constructed by Brandi Chiapelli and Dr. James
McCarter (bchiapel@watson.wustl.edu & jmcCarte@watson.wustl.edu) at
Washington University, St. Louis. DNA Sequencing by: Washington
                                                                                                                                                                                                                                                                                     University Genome Sequencing Center St. Louis
                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: McCarter JP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The Washington Univ. Nematode EST Project, 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                               ative full length read
vector to vector length is 95
vector to overtor length is 95
primer: -40RP from Gibco.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="IMAGE:2019185"
/clone_lib="NCI_CGAP_Brn25"
/tissue_type="anaplastic oligodendroglioma"
/lab_host="DH10B"
                                                 /clone_lib="Parastrongyloides trichosuri FL pAMP1 v1 Chiapelli McCarter"
                                                                                                     /organism="Parastrongyloides trichosuri"
/db_xref="taxon:131310"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.6%;
                           _stage="Free_Living"

 Mismatches

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δÃ

Db QΥ

Locus

QУ В

В

RESULT 8 BI322285

KEYWORDS

COMMENT

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BASE COUNT
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AA154655/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE
AUTHORS
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                                                                                                                                                                                                                                                                                   source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 0.6%;
Local Similarity 62.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AA154655
100 bp mRNA linear EST 19-FEB-1997 mg61c08.rl Soares_thymus_NbMT Mus musculus cDNA clone IMAGE:583214
5' similar to SW:KCC4_MOUSE P08414 CALCIUM/CALMODULIN DEPENDENT PROTEIN KINASE TYPE IV CATALYTIC CHAIN; mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Marra M/Mouse EST Project
Washbu HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The WashU-HHMI Mouse EST Project Unpublished (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 100)
Marra,M., Hiller,L., Allen,M., Bowles,M., Dietrich,N., Dubuque,T.,
Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M.,
Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B.,
Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA154655.1 GI:1726483
                                                                                                                                                                                                                                                                                                     Trace considered overall poor quality possible reversed clone: similarity or seq primer: -28M13 rev2 from Amersham High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                            MGI:357862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: mouseest@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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314 286 1810
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/note="Vector: pT773D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; 1st strand cDNA
                                                                                                                                                                          /db_xref="taxon:10090
/clone="IMAGE:583214"
                                               /tissue_type="Thymus"
/dev_stage="4 weeks"
/lab_host="DH10B"
                                                                                                                          /clone_lib="Soares_thymus_2NbMT"
/sex="male"
                                                                                                                                                                                                                                /strain="C57BL/6J"
                                                                                                                                                                                                                                                        /organism="Mus musculus"

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 27.8; DB 13
Pred. No. 2.7e+04;
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                                                                                                                                                                                                                                                                                                                                                                                         on wrong strand
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SOURCE

COMMENT

TITLE

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т85845/с
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3977 ATTCAAAGCAGATCACCATTGCGATCAACGTCCTCCATGAGCATGGCATAGTCCACCGTG 4036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4037 ACATTAAAGGTGCCAATATCTTCCTTA 4063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27 ATCTAAAAGCTAGCAATATTCTCTATA 1
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Millianson, A., Wolldmann, P. and Wilson, R.

Millianson, A., Wolldmann, P. and Wilson, R.
                                                                                                                                                                                                                                                                                                                                                                   Email: est@watson.wustl.edu
Insert Size: 715
High quality sequence starts: 1 High quality sequence stops: 1
Source: IMAGE Consortlum, LLNL This clone is available royalty-free
through LLNL; contact the IMAGE Consortlum (info@image.llnl.gov)
for further information. Trace considered overall poor quality
Insert Length: 715
Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                 High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                            Seq primer: M13RP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ydfoldio.rl Soares fetal liver spleen INFLS Home sapiens cDNA clone IMAGE:1127723 5' similar to gb:x56468_rnal 14-3-3 PROTEIN THETA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T85845.1 GI:714197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HUMAN);, mRNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 0.6%;
Similarity 57.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24 a
               double-stranded cDNA was ligated to Eco RI adaptors
                                                                                                                                                                        /clone_lib="Soares fetal liver spleen lNFLS"
                                                                                                                                                             /sex≖"male"
                                                                                                                                                                                                      /db_xref="taxon:9606"
/clone="IMAGE:112723"
                                                                                                                                                                                                                                               /db_xref="GDB:468340"
                                                                                                                                                                                                                                                                      /organism≔"Homo sapiens"
                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3']; double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT773 vector. RNA provided by Dr. Bertrand Jordan. Library went through two rounds of normalization, and was constructed by Bento Soares and M.Fatima Bonaldo."
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                                                                                                                                                                                                                                                                                                                                                                                                                            www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 177 Std Error: 0.00
Seq primer: -40m13 fwd. ET from Amersham.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cDNA Library Preparation: David B. Krizman, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R. Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Monica Brown, M.D., Elise Kohn, M.D., Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            qi76e03.xl NCI_CGAP_Ov26 Homo sapiens cDNA clone IMAGE:1862428 3',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 98)
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                                                                                                                                                    /note="Organ: ovary; Vector: pampl; mRNA made from papillary serous ovarian carcinoma, cDNA made by oligo-dr priming. Directionally cloned. Size-selected on agarose
                                                                                                          non-amplified.
28 c
                                                                                                                                          gel, average insert size 500 bp.
                                                                                                                                                                                                                                                /tissue_type="papillary serous carcinoma"
/dev_stage="adult"
                                                                                                                                                                                                                              /lab_host="DH10B"
                                                                                                                                                                                                                                                                                            /sex="female"
                                                                                                                                                                                                                                                                                                       /clone_lib="NCI_CGAP_Ov26"
                                                                                                                                                                                                                                                                                                                                /clone="IMAGE:1862428"
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/db_xref="taxon:9606"
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a 20 c 20 g 33 t 10 others
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88 bp mRNA linear EST 25-APR-:
AU254327 3'-directed mouse cDNA library Mus musculus cDNA clone
BEDD0001610 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: kkato@bs.aist-nara.ac.jp,
URL:http://love2.aist-nara.ac.jp/BED/index.html.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8916-5 Takayama, INTEL: 81-743-72-5581
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Kikuya Kato
Graduate School of Biological Sciences
Nara Institute of Science and Technology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MUS MUSCUIUS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                        Arabidopsis thaliana T-DNA flanking sequence GK-203B07-014506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nouse mouse
                                                                                                                                                                                                        genomic survey sequence
AL760836
A pipeline for automated high-throughput generation of FSTs (flanking sequence tags) from Arabidopsis thaliana T-DNA
                                                                                    Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                      Arabidopsis thaliana
                                                                                                                                                                                                                                                               AL760836
                                    and Weisshaar, B.
                                                    Strizhov, N., Li, Y., Rosso, M., Viehoever, P., Dekker, K., Saedler, H
                                                                                                                                                                                          AL760836.1 GI:21500827
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/clone_lib="3'-directed mouse cDNA library"
/tissue_type="brain"
/note="Vector: pGEM-T-easy"
15 c 6 g 18 t
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/db_xref="taxon:10090"
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so, M., Li, Y., S
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Вb οy Вр QΥ

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4900 GCAAGCTGCAGGTTTGTAATGCAAAAGGCTGATTACTGAAATTTAAGAAAAAGGTTCTTT 4959
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http://www.mpiz.koeln.mpg.de/GABI-Kat/.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AV960997 Nori Satoh unpublished cDNA library, cleavage stage embryo Ciona intestinalis cDNA clone cicl14j15 5', mRNA sequence.

AV960997
Kyoto University Sakyo-Ru, Kyoto 606-8502, Japan Tel: 81-75-753-4081 Fax: 81-75-705-1113
                                                                                                                                                                                                                                                                                                                                                                      Ciona intestinalis.
                                                                                                                       Contact: Nori Satoh
Department of Zoology
                                                                                                                                                                                                              Satoh, N., Satou, Y., Kohara, Y. and Shin-i, T. Expressed genes in Ciona intestinalis
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                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
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/note="PCR was performed on DNA from Arabidopsis thaliana
plants ("1) which were transformed with the T-DNA from
vector pAc161. The lines contain one or more T-DNA
insertions. The DNA fragment(s) resulting from the PCR
were directly sequenced to determine the genomic sequence
flanking the insertion. Sequences displaying significant
similarity to the A. thalaina nuclear genome sequence were
processed for submission. T-DNA derived sequences were
removed"
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Pred. No. 5e+04;
Prematches 22;
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C53310
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                                                                  131 CTAATTTGAAAGAAAAATGAATGCACCAAATCAGCCTCCACATAAAGACACTGGAAAAA 190
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                191 CAGTGGAGAATGTGGAAGAATACAGCTATAAGCA 224
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                                                                                                                                           Similarity
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National Institute of Genetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kohara,Y., Motohashi,T., Tabara,H., Watanabe,H., Sugimoto,A.,M., Miyata,A. and Nishigaki,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Caenorhabditis elegans.
Caenorhabditis elegans
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C53310 Yuji Kohara unpublished cDNA Caenorhabditis elegans cDNA clone yk259b3 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             40;
                                                                                                                                                                                                                                                                                                                                                                                     Email: ykohara@lab.nig.ac.jp.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 81-559-81-6854
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yata 1111, Mishima, Shizuoka 411,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Yuji Kohara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                            Fax: 81-559-81-6855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Expression map of the C.elegans genome
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: satoh@ascidian.zool.kyoto-u.ac.jp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rhabditidae; Peloderinae; Caenorhabditis.
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                                                                                                                           Conservative
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                                                                                                                                                                                                              /dev_stage="varied"
7 c 18 g
                                                                                                                                                                                                                                       /clone_lib="Yuji Kohara unpublished cDNA"
/sex="hermaphrodite, male"
/tissue_type="whole animal"
                                                                                                                                                                                                                                                                                                  /organism="Caenorhabditis elegans"
/strain="CB1489 him-8(e1489)"
/db_xref="rtaxon:6239"
/clone="yk259b3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="whole animal"
/dev_stage="cleavage stage embryo"
/note="Vector: pBluescript SK"
10 c 16 g 35 t
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stage embryo"
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/db_xref="taxon:7719"
/clone="cicl14j15"
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Pred. No. 5.1e+04;
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                                                                                                                  Gaps
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                                            SOURCE
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TITLE
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                       Mus musculus
                                                                                                            mRNA sequence.
     Eukaryota; Metazoa; Chordata;
                                                                             BF101686.1 GI:10884212
                                                                                               BF101686
                                                                                                                             601753418F1 NCI_CGAP_Mam1
                                                                                                                                                 BF101686
                                          house mouse.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ogapbs-r@mail.nih.gov
Tissue Procurement: Jeffrey Green M.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalia; Eutheria; Rodentia;
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                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /sex="female, virgin"
/tissue_type="infiltrating ductal carcinoma"
/dev_stage="5 months"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone_lib="NCI_CGAP_Mam6"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /strain="FVB/N"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Mus musculus"
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Pred. No. 5.4e+04;
                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                           82 bp
Craniata; Vertebrata; Euteleostomi;
                                                                                                                           musculus cDNA clone
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                                                                                                                                             mRNA
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IMAGE:3980885 5',
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COMMENT
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                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM
                                                                                                                                                                                        TITLE
                                                                                                                                                                                                                                                                                                             AUTHORS
                                                                                                                                                                    JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4982 GGAAA 4986
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
rissue Procurement: Gilbert Smith, Ph.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LINL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llni.gov
plate: LLAM9177 row: d column: 06
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Plate: LLAM9177 row: d column: 06 High quality sequence stop: 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1999)
                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 87)

Marra, M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dubuque, T., Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M., Geisel, S., Kucaba, T., Tacy, M., Tan, F., Underwood, K., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B., Steptoe, M., Tan, F., Coarse B., Wilson B., and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   87 bp mRNA linear EST 21-MAR-1: mz90a03.r1 Soares mouse lymph node NbMLN Mus musculus cDNA clone IMAGE:720652 5', mRNA sequence. aA267457
                                                                                                                                           Unpublished (1996)
Contact: Marra M/Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                     Washington University School of MedicineP
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                        The WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA267457.1 GI:1904190
                      Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                 Waterston, R
                                                                                                                                                                                                                                  Theising, B., Wylie, T., Lennon, G., Soares, B., Wilson, R. and
                                                                                                                                                                                                                                                                                                                                                                                                                             house mouse
Email: mouseest@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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/organism="Mus musculus"
/strain="FVBN"
/db_xref="taxon:10090"
/clone="IthAGE:3980885"
/clone_lib="NCI_GAP_Maml"
/clone_lib="NCI_GAP_Maml"
/tissue_type="tumor, biopsy sample"
/tissue_type="tumor, biopsy sample"
/dev_stage="10 months, virgin"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note-*Organ: mammary; Vector: pCMV-SPORT6; Site_1: Sall; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT. Library constructed by Life Technologies. Investigator providing samples: Glibert Smith, NIH*

a 36 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.5%; Score 26.6; DB 12; Length 82; 63.1%; Pred. No. 5.4e+04;

 Mismatches

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AUTHORS
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                                         FEATURES
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seg primer:
     cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
cDNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.lln.govy/bbrp/image/image.html
Insert Length: 820 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 87.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oq38f11.s1 NCI_CGAP_Kid5 Homo sapiens cDNA clone IMAGE:1588653 3',
                                                                                                                                                                                                                                                                       Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1. (bases 1 to 95)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3/]; double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not and Eco RI sites of the modified pT7T3 vector. RNA provided by Dr. Bertrand Jordan. Library constructed and normalized by Bento Soares and M.Fatima Bonaldo." a 10 g 12 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="IMAGE:720652"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="Organ: lymph node; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; lst strand cDNA was primed with a Not I - oligo(dT) primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /tissue_type="lymph node"
/dev_stage="4 weeks"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone_lib="Soares mouse lymph node NbMLN"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.5%; Score 26.4; DB 9; Length 87; 61.8%; Pred. No. 6.3e+04; ative 0; Mismatches 26; Indels
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95
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BASE COUNT
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R93104
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                                                                                                                                                                                                                                                                                                          Seq primer: sk
                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: M.A. Saber, H. Hamied, W.M. El Yassaki, M. Romeih, H. Ahmed, M. Mohareb, I. El Dabaa, S. Mamdouh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Saber,M.A., Hamied,H., ElYassaki,W.M., Romeih,M., Ahmed,H., Mohareb,M., ElDabaa,I. and Mamdouh,S.
                                                                                                                                                                                                                                                                                                                         Email: M-Saber@FRCU.EUN.EG
                                                                                                                                                                                                                                                                                                                                                                      Tel: 202 3128276
                                                                                                                                                                                                                                                                                                                                                                               Theodor Bilharz Research Institute Imbaba, P.O.Box 12411, Giza, Egypt
                                                                                                                                                                                                                                                                                                                                                                                                                                TBRI Biochemistry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Schistosoma mansoni cDNAs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Strigeidida; Schistosomatoidea; Schistosomatidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Schistosoma mansoni,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
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          Note-"Vector: pBluescript II SK+; Site_1: EcoR1; Site_2: XhO1; mRNA was purified from adult couples of S. mansoni. CDNA was constructed and cloned simultaneously using vector priming with the pBluescript II SK+ vector. cDNA was directionally synthesized from the EcoR1 site in the vector to the XhO1 site."
                                                                                                                                                                                                                                                                                                                                              3121167
                                                                                                                                     /Clone_lib="S. mansoni cDNA"
/lab_host="E. coli XL Bluel"
                                                                                                                                                                                                   /db_xref="taxon:6183"
                                                                                                                                                                                                                        /strain="Egyptian"
                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                /clone-"SMTBADAMS0038SK"
                                                                                                                                                                                                                                          /organism="Schistosoma mansoni"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      constructed by Bento Soares and M. Fatima Bonaldo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization. Library
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/clone_lib="NCI_CGAP_Kid5"
/tissue_type="2 pooled tumors (clear cell type)"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1588653"
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Pred. No. 6.5e+04;
39 t
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SOURCE

KEYWORDS VERSION В ρy 밁

ORIGIN

·ORIGIN

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498 AAAGCTTACCTCAGTCTCAAAGAAAAAAGACAGGGAGCAAAGAAGAAAAATAC 554

Matches

Conservative

0;

Mismatches

Indels

0;

Gaps

FEATURES

COMMENT

TITLE JOURNAL

AUTHORS

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ORIGIN
                                                                                               BASE COUNT
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R40354/c
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Best Local Similarity 59.2%;
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1 Similarity
37; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                         High quality sequence stop: 48.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         High quality sequence stops: 61 Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.

Insert Length: 2347 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 78)

Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rikkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and Wilson, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seq primer: Promega -21m13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: est@watson.wustl.edu
Insert Size: 2347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ,R., Williamson,A., Wohldmar
The WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               yf81d05.sl Soares infant brain INIB Homo sapiens cDNA clone IMAGE:29017 3', mRNA sequence.
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                                                                             /dev_stage="73 days post natal"
/lab_host="DH10B (ampicillin resistant)"
/note="Organ: whole brain; Vector: Lafinid BA; Site_1: Not
                                                                                                                                                                                                                                           I; Site_2: Hind III;
                                                                                                                                                                                                                                                                                                                                                             /db_xref="taxon:9606"
/clone="IMAGE:29017"
                                                                                                                                                                                                                                                                                                                           /sex="female"
                                                                                                                                                                                                                                                                                                                                     /clone_lib="Soares infant brain lNIB"
                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="GDB:401364"
            0.5%;
            Score 26.2; DB 14
Pred. No. 6.9e+04;
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                            DB 14; Length 78;
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AV551027 Arabidopsis thaliana roots Col-
cDNA clone RZ120b09R 5', mRNA sequence.
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Arabidopsis thaliana
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DNA Res. 7, 175-180 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The First Laboratory for Plant Gene Research
Kazusa DNA Research Institute
Yana 1532-3, Kisarazu, Chiba 292-0812, Japan
Email: asamizu@kazusa.or.jp, URL:http://www.kazusa.or.jp/en/plant/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AV551027.1 GI:8722440
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                                                                                                                                                                                                                                                                                                                                                                               T62011 93 bp mRNA linear EST 14-FEE yb97b12.rl Stratagene lung (#937210) Homo sapiens cDNA clone IMAGE.79103 5' similar to gb:M64716 40S RIBOSOMAL PROTEIN S25 (HUMAN); mRNA sequence.
Hiller, L. Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins Chissoe, S., Dietrich, N., Dubuque, T., Favello, A., Gish, W., Hawkins C., M., Haltman, M., Kucaba, T., Lacy, M., Le, N., Mardis, E., Moore, M.B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., B., Morris, M.B., Tan, F., Thierry-Meg, J., Trevaskis, E., Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M. Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996)
                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 93)
                                                                                                                                                                                                                                                                                                                                        T62011.1 GI:665254
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/strain="Columbia"
/db_xref="rtaxon:3702"
/clone="RZ120b09R"
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/note="Vector: pBluescriptII SK-; Site_1: EcoRI;
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Pred. No. 7.3e+04;
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source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. I Tel: 314 286 1800 Fax: 314 286 1810
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Contact: Wilson RK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High qallty sequence stops: 62
Source: IMAGE Consortium, LLNL; contact the
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High quality sequence stop: 62.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                     B1858071 NTH_MGC_87 Homo sapiens cDNA clone IMAGE:5393853 5'
                                                   Email: cgapbs-remail.nih.gov

Email: cgapbs-remail.nih.gov

Tissue procurement: DCTD/DTP

Tissue procurement: DCTD/DTP

CDNA Library Preparation: Life Technologies, Inc.

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

http://image.lnl.gov

Plate: LLAM12003 row: m column: 22
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1 (bases 1 to 78)

NIH-MCC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                         mRNA sequence.
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                                                                                                                                                                                                                                    Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                      BI858071.1 GI:15998818
                                  High quality sequence stop:
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/clone="IMAGE:79103"
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/db_xref="GDB:482720"
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    Location/Qualifiers
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56.3%; pred. No. .
50; Mismatches
                                                                                                                                                                                                                       Strausberg, Ph.D.
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Pred. No. 7.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     78 bp
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                                                                                                                                                                                                                                                                                                              Seq primer: M13univ Class: shotgun.
                                                                                                                                                                                                                                                                                                                                                             Email: Michele.morgante@usa.dupont.com
Sequences were trimmed to include only high quality bases; forw.
and reverse reads were assembled when significant overlaps were
                                                                                                                                                                                                                                                                                                                                                                                                                                        Dupont Genomics
PO Box 6104, Newark, DE 19714-6104, USA
Tel: 302 631 2638
Fax: 302 631 2607
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genome Res. 11 (10), 1660-1676 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 85) Meyers, B.C., Tingey, S.V. and Morgante, M. Abundance, distribution and transcriptional activity of repetitive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACC clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Suite 200
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G-10g24.f Maize Random Small-insert Genomic Library Zea mays
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                     cloned into pCR-Script.
                             /note="Vector: pCR-Script; Total genomic DNA was nebulized
; ends were polished with Pfu polymerase and the fragments
                                                                                                                     /sex="hermaphrodite"
/tissue_type="leaf"
                                                                              /dev_stage="seedling"
                                                                                                      /cell_type="Young leaf"
                                                                                                                                                       /clone_lib="Maize Random Small-insert Genomic Library"
                                                                                                                                                                                       /db_xref="taxon:4577"
/clone="G-10g24"
                                                                                                                                                                                                                                 /strain="B73"
                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                              /organism="Zea mays"
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/lab_host="DH10B (phage-resistant)"
/note="Organ: breast; Vector: pcMV-SPORT6; Site_1: NotI;
Site_2: SalI; Cloned unidirectionally: ollgo-dT primed.
Average insert size 1.383 kb. Library enriched for full-length clones and constructed by Life Technologies.
Note: this is a NIH_MGC Library."
a 19 c 23 g 17 t
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/clone_lib="NIH_MGC_87"
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                        TACTATTAATATACAAATGGTGAAGTGTGACAAGTTTGATATACTGTAAGCCATACTTTC 82
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html Insert Length: 2310 Std Error: 0.00 Seg primer: -40ml3 fwd. Er from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               High quality sequence stop: 51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                              /note="vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from 3 pooled germ cell tumors, and was then primed with a Not I oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                  /tissue_type="pooled germ cell tumors"
                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="IMAGE:1469253"
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57.3%; Pred. No. 8.1e+04;
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Pred. No. 8.2e+04;
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AI251245
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Arabidopsis thaliana T-DNA flanking sequence GK-119A08-012517,
genomic survey sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          genomic survey sequence. AL757179
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Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (17-JUN 2002) Weisshaar B., Max-Planck-Institut fuer Submitted (17-JUN 2002) Weisshaar B., Max-Planck-Institut fuer Zuechtungsforschung, Carl-von-Linne-Weg 10, Koeln, 50829, Germany This sequence is recovered from the left border of the T-DNA. It indicates an insertion within the locus defined by clone MWF20. The sequences are generated at the MPI for Plant Breeding Research in the context of the GABI-Kat project. GABI-Kat is part of the Germa Plant Genomics program designated 'GABI'. Information on line availability can be found at: http://www.mpiz-koeln.mpg.de/GABI-Kat/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rosso,M., Strizhov,N., Li,Y., Reiss,B., Dekker,K. and Weisshaar,B. A new Arabidopsis thaliana T-DNA mutagenised population (GABI-Kat) for flanking sequence tag based reverse genetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A pipeline for automated high-throughput generation of FSTs (flanking sequence tags) from Arabidopsis thatiana T-DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone_lib="Arabidopsis thaliana T-DNA insertion lines"
/note="pcr was performed on DNA from Arabidopsis thaliana
/note="pcr was performed on DNA from Arabidopsis thaliana
/note="pcr was performed on DNA from Arabidopsis thaliana
/note="pcr was performed on DNA from DNA from
plants (T1) which were transformed with the T-DNA from
vector pAC161. The lines contain one or more T-DNA
insertions. The DNA fragment(s) resulting from the PCR
insertions. The DNA fragment(s) resulting from the PCR
insertions. Sequences displaying significant
similarity to the A. thalaina nuclear genome sequence were
processed for submission. T-DNA derived sequences were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Li,Y., Strizhov,N. and Weisshaar,B.
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/strain="Columbia_0"
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                                                                                                                                                                                                                                                                                                                   0.5%; Score 26; DB 17; Length 88; 65.5%; Pred. No. 8.2e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
unknown library type
Seq primer: -40UP from Gibco.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mRNA sequence.
AI251245
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AI251245.1 GI:3847774
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                        A1035450 78 bp mKNA linear EST 26-JUI ub47b05.rl Soares_mammary_gland_NbMMG Mus musculus cDNA clone IMAGE.1380849 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.5%;
Similarity 57.3%;
                                                                                                                                                                                            Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 78)

1 (bases 1 to 78)

Marra, M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dubuque, T., Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M., Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B., Theising, B., Wylie, T., Lennon, G., Soares, B., Wilson, R. and
                                                                                                                                                                                                                                                                                                                                                                                                         AI035450.1 GI:3259160
                                                    WashU-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                  Contact: Marra M/Mouse EST Project
                                                                                                                                   Unpublished (1996)
                                                                                                                                                             The WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                     house mouse
Email: mouseest@watson.wustl.edu
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314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1985542"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="papillary serous carcinoma"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="NCI_CGAP_Ov32"
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Pred. No. 8.
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FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 TGTAAACACGGCACCACAGAGTGTTTTCAGAATGGCTTCCTTTGGGTGGAAGAGGAGAAT 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
            www-bio.lln1 gov/bbrp/image/image.html
Seq primer: -40ml3 fwd. ET from Amersham.
                                                 found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R. Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                   NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
                                                                                                                                                                                                                                               Unpublished (1997)
                                                                                                                                                                                                                                                                                                                       Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 83)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA915897 83 bp mRNA linear EST 14-APR-1998 oh86b08.s1 NCI_CGAP_CO8 HOMO Sapiens cDNA clone IMAGE:1473879 3' similar to gb:V00567 BETA-2-MICROGLOBULIN PRECURSOR (HUMAN);, mRNA
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                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA915897.1 GI:3055289
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MGI:903317
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="Organ: mammary gland; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: EcoRI; 1st strand cDNA was primed with a Not I - oligo(dT) primer [5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /db_xref="taxon:10090"
/clone="IMAGE:1380849"
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/dev_stage="4 weeks"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone_lib="Soares_mammary_gland_NbMMG"
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Pred. No. 8.8e+04;
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                                                                                                                                                                                                                                                                                                                   Tissue Procurement: Dennis Sgroi, M.D., Kristina Cole, M.D., Ph.D. Student, Michael R. Emmert-Buck, M.D., Ph.D. CDNA Library Preparation: David B. Krizman, Ph.D. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 164 Std Error: 0.00
                                                                                                                                                                                                                                                                                                      Insert Length: 164 Std Erro
Seq primer: -40UP from Gibco.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AI340694 91 bp mRNA linear EST 18-MAR-19 tb53C05.x1 NCI_CGAP_Br15 Homo sapiens cDNA clone IMAGE:2058056 37 similar to gb:X06989_rna1 ALZHEIMER'S DISEASE AMYLOID A4 PROTEIN
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adenocarcinoma tissue, cDNA made by oligo-dr
               /lab_host="DH10B"
/note="Organ: breast; Vector: pAMP1; mRNA made from breast
                                                                                /tissue_type="adenocarcinoma"
/dev_stage="adult"
                                                                                                                                 /sex="female'
                                                                                                                                                                               /clone="IMAGE:2058056"
                                                                                                                                                                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                        /clone_lib="NCI_CGAP_Br15"
                                                                                                                                                                                                                                                                               Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:9606"
/clone="IMAGE:1473879"
/clone_lib="NGI_CGAP_CG8"
/tissue_type="adenocarcinoma"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
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mc67e06.x1 Soares mouse embryo NbME13.5 14.5 Mus musculus cDNA
clone IMAGE:353602 3' similar to SW:RPB2_HUMAN P30876 DNA-DIRECTED
RNA POLYMERASE II 140 KD POLYPEPTIDE ;, mRNA sequence.
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Marra, M., Hillier, L., Kucaba, T., Martin, J., Beck, C., Wylie, T., Underwood, K., Steptoe, M., Theising, B., Allen, M., Bowers, Y., Person, B., Swaller, T., Gibbons, M., Pape, D., Harvey, N., Schurk, R., Ritter, E., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R., Waterston, R. and Wilson, R. R. 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (1999)
Contact: Marra M/WashU-NCI Mouse EST Project 1999
Contact: Marra M/WashU-NCI Mouse EST Project 1999
Washington University School of Medicine
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108, USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. This clone was previously sequenced on the 5' end only, this new data is from the 3' end.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: mouseest@watson.wustl.edu
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ilarity 58.4%;
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/note="Vector: pT/T3D-pac (Pharmacia) with a modified /note="Vector: pT/T3D-pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Ist Strand CDNA polylinker; Not I - oligo(dT) primer [5', primer RI] of the primer RI] oligo(dT) primer RI] of the primer RI] on equal amounts of mRNA from 2 13.5dpc and 2 7 3/1, on equal amounts of mRNA from 2 13.5dpc and 2 14.5dpc embryos (total RNA provided by Minoru Ko, Wayne State Univ., from 2 1; double-stranded cDNA was ligated to Broat adaptors (Pharmacia), digested with Not I and Broat RI sites of the modified cloned into the Not I and Broat RI sites of the modified
                                                                                                                                                                                                                                                                                                                                                                                                                                   /tissue_type="embryo"
/dev_stage="13.5-14.50
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/clone="IMAGE:353602"
/clone_lib="Soares mouse embryo NbME13.5 14.5"
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/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                      /lab_host="DH10B"
/note="Vertical fetus"
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                                                                                                                 AI142956
                                                                                                                                   RESULT 34
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 505 ACCTCAGTCTCAAAGAAAAAAGACAGGGAGCAAAGAAGAAGAAAATACGTC 557
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                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Taylor,R., Ashurst,J.L., Croning,M.D.R., Zorn,A.M. and Rogers,J. sanger Xenopus tropicalis EST project 2002 (npublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AL794139 AL794139 XGC-neurula Silurana tropicalis cDNA clone TNeul16m13 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AL794139.1 GI:21579843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mRNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Taylor R
Sanger Centre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    western clawed frog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequencing primer: PlCSP6
This sequence is from a X
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sanger Xenopus tropicalis EST project 2001 TROPICALIS_SEQUENCE_ID: TNeullim13.plcSP6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  xenopodinae; Silurana.
                                                                                                                                                                                                                                                                    1 Similarity
36; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       constructed by Aaron M. Zorn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: trop@sanger.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HINKEOU'
                                 AI142956 73 bp mRNA linear EST 25-SEP-1 oz58d01.x1 Soares_senescent_fibroblasts_NbHSF Homo sapiens cDNA clone IMAGE:1679521 3' similar to SW:SR54_HUMAN P13624 SIGNAL RECOGNITION PARTICLE 54 KD PROTEIN ;, mRNA sequence.
    AI142956.1 GI:3659315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  M.Fatima Bonaldo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cambridgeshire, CB10 1SA, UK
                                                                                                                                                                                                                                                                                                                                                                   /dev_stage="neurula"
/dev_stage="neurula"
/lab_host="Escherichia coli DH10B"
/lab_host="secherichia coli DH10B"
/lab_host="secherichia coli DH10B"
/note="Vector: pCS107; Site_1: EcoRI; Site_2: NotI; cDNA
was oligo dT primed from 5ug of poly A+ RNA from neurula.
was oligo dT primed from 5ug of poly A+ RNA from neurula.
EcoRI NotI cut cDNA was then ligated into pCS107 with
EcoRI at the 5' end and NotI at the 3' end."

ECORI at the 5' end and NotI at the 3' end."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Silurana tropicalis"
/db_xref="taxon:8364"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone_lib="XGC-neurula"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone="TNeu116m13"
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                                                                                                                                                                                                                                                                                          Score 25.8; DB 9;
Pred. No. 9.7e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a Xenopus Gene Collection (XGC) library
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10 ATTCCATCATGCCTTTCATGTTGCCAGCAGCACCCTGGTGAAACTGCCTTATCAGTGACT 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
Email: Cyapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Insert Length: 1214 Std Error: 0.00
                                                                                          Contact: Robert Strausberg, Ph.D.
                                                                                                                  Unpublished (1997)
                                                                                                                                        NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                            OZ44c01.x1 Soares_NhHMPu_S1 Homo sapiens cDNA clone IMAGE:1678176
                                                                                                                                                                                                                                                                                                                                                                                  AI086378
                                                                                                                                                                                                                                                    HOMO sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  40;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
Seq primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP).
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1 (bases 1 to 73)
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                                                                                                                                                                                                                                                                                                                                      , mRNA sequence.
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                                                                                                                                      Gene Index
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library went through one round of normalization to a Cot = 5. Library constructed by Bento Soares and M.Fatima Bonaldo."
                                                                                                                                                                                            1 to 85)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="IMAGE:1679521"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            62.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.5%; Score 25.6; DB 9; 62.5%; Pred. No. 9.8e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                        Kazusa DNA Research Institute
Yana 1532-3, Kisarazu, Chiba 292-0812, Japan
Email: asamizu@kazusa.or.jp. URL:http://www.kazusa.or.jp/en/plant/.
                                                                                                                                                                                                                                                                  The First Laboratory for Plant Gene Research
                                                                                                                                                                                                                                                                                                                                                         Asamizu.E., Nakamura,Y., Sato,S. and Tabata,S. A large scale analysis of CDNA in Arabidopsis thaliana: Generation of 12,028 non-redundant expressed sequence tags from normalized and size-selected CDNA libraries
                                                                                                                                                                                                                                                                                                  Contact: Erika Asamizu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AV532477 Arabidopsis thaliana flower buds Columbus thaliana cDNA clone FB042h0lf 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                           20363093
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 (bases 1 to 86)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17 a
/clone_lib="Arabidopsis thaliana flower buds Columbia"
/tissue_type="flower buds"
/note="Vector: pBluescriptII SK-; Site_1: EcoRI; Site_2:
                                                                                                                                                                                                                                                                                                                                              7, 175-180 (2000
                                                                                  /clone="FB042h01F"
                                                                                                   /strain="Columbia" /db_xref="taxon:3702"
                                                                                                                                   /organism="Arabidopsis thaliana"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="Organ: mixed (see below); Vector: pT773D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Eggal amounts of plasmid DNA from three normalized libraries (melanocyte 2NbHM, pregnant uterus NbHPU, and fetal heart NbHHJ9W) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR amplified CDNAs from pools of consisted of I.M.A.G.E. clones 260232-265223, 340448R-74470 and 404408-200425.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 340488-345479, and 484488-489479."

6 c 8 g 54 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone_lib="Soares_NhHMPu_S1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="IMAGE:1678176"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
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57.5%;
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Pred. No. 1e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              82 AGCA 85
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zuechtungsforschung, Carl-von-Linne-Weg 10, Koeln, 50829, Germany This sequence is recovered from the left border of the T-NNA. It indicates an insertion within the locus defined by clone F4F7. The sequences are generated at the MPI for Plant Breeding Research in the context of the GABI-Kat project. GABI-Kat is part of the German Plant Genomics program designated 'GABI'. Information on line availability can be found at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 (bases 1 to 0/)
11,Y., Strizhov,N., Rosso,M. and Weisshaar,B.

Li,Y., Strizhov,N., Rosso,M. and Weisshaar,B.

Direct Submission
Submitted (17-JUN-2002) Weisshaar B., Max-Planck-Institut fuer
Submitted (17-JUN-2002) Weisshaar B., Max-Planck-Institut fuer
Transphing Carl-von-Linne-Weg 10, Koeln, 50829, Germany
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A pipeline for automated high-throughput generation of FSTs (flanking sequence tags) from Arabidopsis thaliana T-DNA transformed lines
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rosso,M., Strizhov,N., Li,Y., Reiss,B., Dekker,K. and Weisshaar,B. A new Arabidopsis thaliana T-DNA mutagenised population (GABI-Kat) for flanking sequence tag based reverse genetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Strizhov, N., Li, Y., Rosso, M., Viehoever, P., Dekker, K., Saedler, H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
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                                                                                                                                                     /clone_lib="Arabidopsis thaliana T-DNA insertion lines" /note="PCR was performed on DNA from Arabidopsis thaliana plants (T1) which were transformed with the T-DNA from vector pACIG1. The lines contain one or more T-DNA insertions. The DNA fragment(s) resulting from the PCR were directly sequenced to determine the genomic sequence flanking the insertion. Sequences displaying significant similarity to the A. thalaina nuclear genome sequence were
                                                                                                                                  processed for submission. T-DNA derived sequences were
                                                                                                                                                                                                                                                                                                                                                                          /clone="GK-059E02-013873"
                                                                                                                                                                                                                                                                                                                                                                                               /strain="Columbia 0"
/db_xref="taxon:3702"
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/strain="Columbia O"
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Score 25.6; DB 17; Length 87;
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VERSION
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                                                                DEFINITION
                                                                                                                                 RESULT 39
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                                                                                                                                                                                                                                      1959 TAGTTTAAAAAATCT 1973
                                                                                                                                                                                                                                                                                                                      1899 GTATTTTGATTACATGAGAAGCTGGATCCAAATGCTACAGCAATTACCTCAAGCATCGCA 1958
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                                                                                                                                                                                               65 TACTTTATCAAATGT 79
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Local Similarity 58.7%;
                                                                                                                                                                                                                                                                              5 GTTTTATGATTTATTTAACTTGTGGAACAAAAATAAACCAGATTAACCACAACCATGCCT 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    34;
                 2M0230B17F Mouse 10kb plasmid UUGC2h
clone UUGC2M0230B17 F, DNA sequence.
A2961698
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
Email: cgapbs -r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 79)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            High quality sequence stop: 67.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      www-bio.llnl.gov/bbrp/image/image.html
Seg primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CDNA Library Preparation: Life Technologies, Inc. CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (1997)
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  AZ961698.1 GI:13832925
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                 AZ961698
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: SalI; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.75 kb. Life Technologies catalog #: 11538-014"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              adenocarcinoma, 7 pooled tumors"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone_lib="NCI_CGAP_Utl"
/tissue_type="well-differentiated endometrial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone="IMAGE:2629258"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /db_xref="taxon:9606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
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Pred. No. 1.1e+05;
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                                                                                                                                                          4114 AAAAACAATGCCCAG 4128
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    D19910
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Insert Length: 10000 Std Error: 0.00
Plate: 0230 row: B column: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2000)
Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                High quality sequence stop: 89.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plasmid inserts
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                            adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

17 c 16 g 28 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114) gblaff129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab_host="E. coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (female) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Biomedical Polymers Research Bldg.,
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/clone="UUGC2M0230B17"
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/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone_lib="Mouse 10kb plasmid UUGC2M library"
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                                                                                                                                                                                                                                                                                                 0;
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Pred. No. 1.2e+05;
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                 4936 TGAAATITAAGAAAAAGGTTCTTTTTTCAATAAATGGTTTATTTTAGGAAA 4986
                                                                                                                                                                                                                                                                                                           source
41 TGAACTCTAAGAAATAAATTGGTTTGGTAATAAATGGCTTCTGGTCAGAAA
                                                                             35;
                                                                                                                                                                                                                                                                                                                                                                                 Contact: Okubo,K., Fukushima,A., Yoshii,J., Niiyama,T., Kojima,Y., Yoshinari,H., Arimoto,J. and Matsubara,K.
Institute for Molecular and Cellular Biology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Okubo, K., Fukushima, A., Yoshii, J., Niiyama, T., Kojima, Y., Yoshinari, H., Arimoto, J. and Matsubara, K.
Gene expression of human promyelocytic cell line HL60 before and after induction of differentiation. A new application of 3'directed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 91)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HUMGS00867 Human promyelocyte Homo sapiens cDNA clone mm0905 3',
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               cDNA sequencing
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                                                                                                                                                                                                                                                                                                                                                                 Osaka University
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                                                                           Conservative
                                                                                                                                                                       32
                                                                                                                                                                     promyelocyte.
                                                                                                                                                                                             /note="Female, adult, cell_line = HL60, cell_type =
promyelocyte."
                                                                                                                                                                                                                                               /db_xref="taxon:9606"
/clone="mm0905"
                                                                                                                                                                                                                           /clone_lib="Human promyelocyte"
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Pred. No. 1.2e+05;
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Search completed: November 11, 2002, 10:46:34 Job time : 4287 secs

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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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Coupled ligation

Human bone marrow Probe #21912 used Thymosin alpha-1 n HIV DNA encoding G Sbg1 exon sequence Human genome-deriv Human foetal liver Probe #18368 for g Human brain expres Human bone marrow Probe #16913 for g Probe #24484 used Human MEKK4 DNA am Human neutrophil c 80%E Bundle peptid Human foetal liver Probe #14054 for 9 Mouse spliced tran Transforming growt cDNA encoding Dros HCV DNA encoding H Human neutrophil c Expression vector
Human foetal liver
Probe #15434 for g
Human brain expres Synthetic storage Synthetic lysine-r Human SNP oligonuc Human SNP oligonuc Mouse spliced tran Sequence encoding Human gene signa Cell death protect Clone 92-2 used in SSP 5.11.11.5 clon Human secreted pro Human secreted pro Reverse PCR primer GFP Leu(CTG)5 forw Human foetal liver Synthetic GFP olig Mercuric Mouse spliced tran Mouse spliced tran Synthetic transcri Probe #20956 used Sbg1 exon sequence Human Probe #14775 for g Human spliced tran Human Probe #19061 used exon sequence gene signatu brain expres insulin deri foetal liver brain expres foetal genome-deriv ion

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    (COMP-) COMPUGEN INC
             28-JUL-2000; 2000US-221607P
02-MAY-2001; 2001US-287724P
                           20-JUL-2001; 2001WO-IB01903
                                                W0200210449-A2
                                                         Homo sapiens
                                                                   splice variant;
                                                                              Human spliced transcript detection oligonucleotide SEQ ID NO:8213.
                                                                                          15-JUL-2002 (first entry)
                                                                                                             ABN35465 standard; DNA; 60 BP
                                                                                                                                                      mouse; rat; splice transcript; detection; RNA transcript
                                                                                                                                                     transcriptome; oligonucleotide library; ss.
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Human bone marrow
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Probe #15216 for g
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Matches Query Match

Local

h 1.2%; Similarity 100.0%;

Conservative

0; Mismatches Score 60; Pred. No.

DB 24; .8e-07; 0

Length 60 Indels

0; Gaps

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CC biological sample, in expression profiling studies, in qualitatively or cypantitatively characterising the corresponding transcriptome, and in transcriptomes. The libraries may also be used as specialised minic transcriptomes. The libraries may also be used as specialised minic constraints to detect transcripts of a sub-transcriptome under a comparticular biological or pathological state, and so allowing the constraints of tissue—and pathological state, and so allowing the constraints of a specific pathological constraints of a specific pathological constraints, to detect developmental specific genes such as those genes constraints and splice variants of a transcriptome of a patient suffering from a particular disorder. ABN27253 to ABN59589 represent coligonucleotide sequences from rats, humans and mice, which are used in the exemplification of the present invention.

CC specification, but was obtained in electronic format directly from WIPO at this winn intimation of the present compared to the printed specification, but was obtained in electronic format directly from WIPO at this winn intimation is patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
Sequence 60 BP; 17 A; 17 C; 10 G; 16 T; 0 other;
                                                   at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes oligonucleotide libraries for detecting messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multiple transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to a set of messenger RNAs transcribed from a given transcription unit of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the genome, which encodes one or more messenger RNA splice variants. The oligonucleotide libraries are useful for detecting mRNAs from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription unit of a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; SEQ ID 8213; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          developmental-specific genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shoshan A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-257383/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wasserman A, Mintz E, Mintz L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Faigler S;
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DE

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KW

KW

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PN

PN

PD

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PR

PR

PA

XX
                                                                                                                                                                                                                                                                   RESULT 2
ABN55582/c
                                                                                                                                                                                                                                                                                                                밁
                              28-JUL-2000;
02-MAY-2001;
                                                                                                                                               Human; mouse; rat; splice transcript; detection; RNA transcript; splice variant; transcriptome; oligonucleotide library; ss.
                                                            20-JUL-2001; 2001WO-IB01903.
                                                                                     07-FEB-2002.
                                                                                                                                                                              Mouse spliced transcript detection oligonucleotide SEQ ID NO:28330.
                                                                                                                                                                                                        15-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                      WO200210449-A2.
                                                                                                                                  Mus musculus.
                                                                                                                                                                                                                                   ABN55582;
                                                                                                                                                                                                                                                     ABN55582 standard; DNA; 65 BP
                             2000US-221607P
2001US-287724P
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(COMP-) COMPUGEN INC

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The present invention describes oligonucleotide libraries for detecting cc messenger RNAs that populate a (sub-)transcriptome, where the cysub-)transcriptome comprises messenger RNAs transcribed from multiple ctranscription units that populate a genome. The library comprises cc several oligonucleotides, each capable of hybridising selectively to a set of messenger RNAs transcribed from a given transcription unit of the genome, which encodes one or more messenger RNAs plice variants. Cc the genome, which encodes one or more messenger RNAs plice variants of cupitodical sample, in expression profilling studies, in qualitatively or quantitatively characterising the corresponding transcriptome, and in cc detecting RNA transcripts and splice variants of human or animal cupitodical suppose the corresponding transcriptome under a particular biological or pathological state, and so allowing the conjugation of tissue and pathology specific genes such as those genes conjugations; to detect developmental specific genes such as those genes conjugations; to detect developmental specific genes; and to detect RNA cc condition; to detect developmental specific genes; and to detect RNA cc and splice variants of a transcriptome of a patient suffering cc particular disorder. ABN2723 to ABN59589 represent coligonucleotide sequences from rats; humans and mice, which are used in cc the exemplification of the present invention.

CC N.B. The sequence data for this patent did not form part of the printed cc at ftp.wipo.int/pub/published_pct_sequences.
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  AAD37143
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription unit of a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-257383/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shoshan A, Wasserman A, Mintz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; SEQ ID 28330; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         developmental-specific genes
                                                                                                                                                Human; MEKK4 modulation; mitogen-activated protein kinase kinase 4; MTK1; MAP3K4; MAP three kinase 1; MAP/ERK kinase 4; MAPKKK4; cytostatic; prophylaxis; immunological; hyperproliferative disorder; cancer; therapy; antisense; inflammatory; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1601 GGGA 1604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 65 BP; 17 A; 18 C; 15 G; 15 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                AAD37143 standard; DNA; 28 BP
                                                                                                                                                                                                                                              Human MEKK4 DNA amplifying forward PCR primer
                                                                                                                                                                                                                                                                                                                            AAD37143;
                                                                                                                                                                                                                                                                                         21-AUG-2002
28-SEP-2001; 2001WO-US30549
                                                                           WO200227033-A1
                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                              5 GAGA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 44.8;
Pred. No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mintz L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.0066;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Faigler S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 65;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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Ър
                                                                                              Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to antisense compounds, compositions and methods for modulating the expression of MEKK4 (also referred as mitogenactivated protein kinase kinase 4; MAP3K4; MAP three kinase 1; MAP/ERK kinase 4; MAPKKK4; MTK1). The antisense oligos are useful for inhibiting the expression of MEKK4 in cells or tissues. They are also useful for treating an animal having a disease or condition associated with MEKK4 such as immunological, inflammatory, hyperproliferative disorder or cancer. Sequences of the invention are also useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. They are also useful in antisense therapy. The present sequence is a PCR primer which is used for amplifying human MEKK4 DNA. This sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compound targeted to nucleic acid encoding mitogen-activated protein kinase 4, useful for treating immunologic discreter. inflammators discreter or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 13; Page 89; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disorder, inflammatory disorder or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-416486/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ward DT, Gaarde WA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-SEP-2000; 2000US-0676436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 28 BP; 9 A; 5 C; 6 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                          07-APR-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT50934;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT50934 standard; cDNA; 78 BP
                                                                                                                                                                                                                                                                                                                                                                                          Mouse p53-recognition clone 1 5' sequence.
Disclosure; Fig 2a; 30pp; English
                                                                                                                                                                                                                                                                                                                                                          p53 binding protein; Mdmx; tumour suppressor; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        784 ACTCCTGGAACAAAGATTGTAGGTTACT 811
                                                                                                                                               (UYLE-) RIJKSUNIV LEIDEN.
                                                                                                                                                                                               13-JUN-1995;
                                                                                                                                                                                                                               13-JUN-1996;
                                                                                                                                                                                                                                                               27-DEC-1996.
                                                                                                                                                                                                                                                                                               W09641875-A1
                                  mouse Mdm2,
                                                 Nucleic acid encoding p53-binding protein - which has homology to
                                                                                 WPI; 1997-065462/06.
                                                                                                               Jochemsen A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 ACTCCTGGAACAAAGATTGTAGGTTACT 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                  for use in cancer research
                                                                                                                 Shvarts A,
                                                                                                                                                                                                 95EP-0201565
                                                                                                                                                                                                                               96WO-NL00239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Monia BP, Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.6%; Score 28; DB 24;
100.0%; Pred. No. 3.8e+02;
                                                                                                                   Van Der Eb AJ;

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 28÷
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AAC19445/c
ID AAC19
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Best Local :
            The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences cDNA sequences in the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. SeTs are derived from mRNAs with intact 5' ends and can therefore be in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design
       expression and secretion vectors
                                                                                                                                                                                                                                                             Claim 1; SEQ ID 23520; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                          obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                             New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for
                                                                                                                                                                                                                                                                                                                                                                                                          Dumas Milne Edwards J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-FEB-2000; 2000EP-0200610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene therapy; chromosome mapping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EP1033401-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human secreted protein 5' EST, SEQ ID NO: 23520.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-OCT-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAC19445 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1182 TGAAGAAGAACAAA 1195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1122 GTATGAGGGTGATGACACAGAAGGAGAATTAAAGGAGTTGGAAAGTAGTACGGATGAGAG 1181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC19445;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 78 BP; 33 A; 9 C; 22 G; 10 T; 4 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The sequences of the 5' region (AAT50934) and 3' region (AAT50935) were detd. for cDNA clone 1, isolated from a mouse 16-day embryo cDNA library using tumour suppressor p53 as probe. The 5' and 3' regions (AAT50936-37) of another clone were also detd. These sequences showed no significant homology to any known sequence submitted to several databases. A third clone (see also AAT50933)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            coded for a p53 binding protein designated Mdmx (AAW10206).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 TGVNGHAGACAAAA 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GTATGAGGTGGAAGAACAGAAGTGGTCATNAAGTCATACCAGAAGAACAGCGATCAGGA 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                         2000-500381/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5' EST;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    43;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0122487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          expressed sequence tag; secreted protein; cDNA isolation;
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                                                                                                                                                                                                                                                                                                                                                                                                        Duclert A,
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                        Giordano J;
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hes 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
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RESULT 6
AAD37145
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
The present invention relates to antisense compounds, compositions and methods for modulating the expression of MEKK4 (also referred as mitogenactivated protein kinase kinase 4; MAPJR4; MAP three kinase 1; MAP/ERK kinase kinase 4; MAPKK4; MAP three kinase 1; MAP/ERK kinase kinase 4; MAPKK4; MAP three kinase 1; MAP/ERK kinase kinase 4; MAPKK4; In cells or tissues or useful for inhibiting the expression of MEKK4 in cells or tissues. They are also useful for treating an animal having a disease or condition associated with MEKK4 such as immunological, inflammatory, hyperproliferative disorder or cancer. Sequences of the invention are also useful for diagnostics, therapeutics, prophylaxis and as research reagents and kits. They are also useful in antisense therapy. The present sequence is a PCR probe which is used for amplifying human MEKK4 DNA. This sequence
                                                                                                                                                                                           Example 13; Page 89; 132pp; English.
                                                                                                                                                                                                                         disorder, inflammatory disorder or cancer
                                                                                                                                                                                                                                    New antisense compound targeted to nucleic acid encoding mitogen-activated protein kinase 4, useful for treating
                                                                                                                                                                                                                                                                                     WPI; 2002-416486/44.
                                                                                                                                                                                                                                                                                                                     Ward DT,
                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                             29-SEP-2000; 2000US-0676436
                                                                                                                                                                                                                                                                                                                                                                                                         28-SEP-2001; 2001WO-US30549
                                                                                                                                                                                                                                                                                                                                                                                                                                           04-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200227033-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; MEKK4 modulation; mitogen-activated protein kinase kinase 4; MTK1; MAP3K4; MAP three kinase 1; MAP/ERK kinase kinase 4; MAPKKK4; cytostatic; prophylaxis; immunological; hyperproliferative disorder; cancer; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antisense; inflammatory; PCR; probe; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human MEKK4 DNA amplifying PCR probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 92 BP; 12 A; 9 C; 15 G; 47 T; 9 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            143 AAAAAATGAATGCACCAAATCAGCCTCCACATAAAGACACTGGAAAAACAGTGGAGAATG 202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          203 TGG 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TGG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  36;
                                                                                                                                                                                                                                                                                                                    Gaarde WA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "FAM labelled 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "TAMRA labelled guanine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag= b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.5%;
                                                                                                                                                                                                                                                                                                                  Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 26.6; DB 21
Pred. No. 2.3e+03;
                                                                                                                                                                                                                                                                                                                Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             thymine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 92;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                      immunologic
                                                                                                                                                                                                                                                                                                                                                                                                              nt.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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RESULT 7
AAQ86400/c
ID AAQ864
XX
                                                                                                                                                                                                           Вb
                                                                          δÃ
                                                                                                      B
                                                                                                                           QΥ
RESULT 8
ABK55302/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                          Matches
                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 26 BP; 5 A; 8 C; 6 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ86400 standard; DNA; 100 BP
                                                                                                                                                                                                                                 AAQ86399 and AAQ86400 are a pair of primers for the PCR amplification of the Asp3 B-chain/Ala21 A-chain human insulin derivative DNA. The two subunits of the derivative are covalently connected via disulphide bonds between Cys residues A7/B7 and A20/B19. The derivative, which may be present as a zinc ion complex, can be used as a fast action treatment for diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-SEP-1993;
02-FEB-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9507931-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ala21 PCR primer #128;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human insulin derivative; A-chain; diabetes; Zinc ion complex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human insulin derivative Ala21 A-chain PCR primer #128.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-NOV-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ86400;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       828 TCTCCAACGCCAGAGGGTCTCATTTG 853
                                                                                                                                                                                                                                                                                                                                                         Acylated insulin deriv. which may be present as a Zinc ion complex - is used to treat diabetes and is rapid acting.
                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1995-131314/17.
                                                                                                                                                                                                                                                                                                                                                                                                                         Andersen AS,
Markussen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-MAR-1995
                                                                          4529 TATTTACATAAAGACTGTGCTGAGAAGCAG 4558
                                                                                                                              Sequence 100 BP; 29 A; 22 C; 23 G; 26 T; 0 other;
                                                                                                                                                                                                                                                                                                                             Example 1; Page 30; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                       Markussen
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NOVO ) NOVO-NORDISK AS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 TCTCCAACGCCAGAGGGTCTCATTTG 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                     37 CAATTGGAAAACTACTGTGCTTAGACGCAG 8
                                                                                                       97 AAGTCTGACGACGCTAAGGGTATCGTTGAACAATGTTGTACTTCTATCTGTTCTTTTGTAC 38
                                                                                                                                                                      ocal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26;
                                                                                                                                                          50;
                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                     Halstrom JB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93DK-0001044.
94US-0190829.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94WO-DK00347.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                      0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 26; DB 24; pred. No. 1.4e+0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Havelund S,
                                                                                                                                                                       Score 26; DB 16; Length 100; Pred. No. 3.6e+03;
                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.4e+03;
tes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Jonassen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                 Db
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RESULT 9
                                                                                                                         AAH48689/c
                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                               of cancer in a patient. (I) and (II) are useful in pharmaceutical compositions, e.g. vaccines, and other compositions for the diagnosis and treatment of colon cancer. A composition comprising a first component selected from physiologically acceptable carriers and immunostimulants, and an antigen-presenting cell expressing (II) is useful for inhibiting development of cancer in a patient. (I) is useful in the design and preparation of ribozyme molecules for inhibiting expression of tumour polypeptides and (I). ABK54531-ABK55464 represent human colon cancer cDNA sequences of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK55302 standard; cDNA; 81 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-AUG-2000; 2000US-223265P.
02-OCT-2000; 2000US-237405P.
20-MAR-2001; 2001US-277495P.
03-JUL-2001; 2001US-302702P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; colon cancer; immunogenic; vaccine; tumour; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human colon cancer-associated cDNA, SEQ ID No 772.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to isolated polynucleotides (I) encoding colon tumour polypeptides (II). (I) is useful for stimulating an immune response in a patient and treating colon cancer in a patient oligonucleotides derived from (I) are useful for determining the presence oligonucleotides derived from (I) are useful for determining the presence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-FEB-2002
                                                                                                                                                                                                                                                                                   3723 CACCTGCATCAGCGTCGACACCGGGGAGCTGATGGCCATGAAAGAGATTCGATTTCAACC 3782
                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 81 BP; 9 A; 22 C; 15 G; 33 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 368; 425pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel polynucleotide encoding colon tumour polypeptides, useful as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-257462/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pyle RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-JUL-2001; 2001WO-US23826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200212280-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     vaccines for treating colon cancers
                                                                                                                                                                                                                             3783 TAATGACCATAAGACTATC 3801
Human G-protein subunit Gbeta3 intron 9 DNA fragment #2.
                                                                         AAH48689;
                                                                                                       AAH48689 standard; DNA; 60 BP
                                      19-OCT-2001 (first entry)
                                                                                                                                                                                                                                                             79 CAGCAGAAGCAGGGTACCCTTTGGGGAGCAGGAGGCCAATAAAAAGTTCAGAGTTCAAAA 20
                                                                                                                                                                                          19 AAAAAAAAAAAGGGCNNCC 1
                                                                                                                                                                                                                                                                                                                                                  ocal Similarity
                                                                                                                                                                                                                                                                                                                                    45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Xu J, Secrist H;
                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                  0.5%;
                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                  Score 25.8; DB 24;
Pred. No. 3.6e+03;
                                                                                                                                                                                                                                                                                                                                        34;
                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                       Length 81;
                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                            Gaps
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0;

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RESULT 10
AAA71594
ID AAA71
XX
AC AAA71
XX
AC AAA71
XX
DF Human
XX
Brain
KW Brain
KW Brain
KW Heart
XX
OS Homo
XX
PN WO20(
XX
PD 03-AI
XX
PR 27-J;
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
     29-JAN-1999;
                                                       27-JAN-2000;
                                                                                                              03-AUG-2000
                                                                                                                                                             W0200045176-A2
                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                              Brain natriuretic
                                                                                                                                                                                                                                                                                                                                     Human brain natriuretic peptide N-terminal primer Pro2hum
                                                                                                                                                                                                                                                                                                                                                                                                       11-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA71594 standard; DNA; 70 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              colitis ulcerosa, transplant rejection, HIV or Hepatitis B or C. This sequence represents a fragment of the human G-protein Gbeta3 subunit intron 9 which is used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G-protein beta3 subunit to predict physiological and pathological pathways in the human body. The invention may be used to detect predisposition for, give a prognosis of, or treat a variety of diseases including high blood pressure, hypertonia, cardiac infarction, coronary disease and other heart or circulation diseases, diabetes, psychiatric diseases, immunological diseases, asthma, psociasis, Crohn's disease, asthma, psychiatric diseases, crohn's disease, asthma, psychiatric diseases, asthma, psychiatric dise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 60 BP; 6 A; 11 C; 6 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel use for sequence variants of the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 3; 14pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Use of sequence variants of the beta3 subunit of human G predict, detect, and determine drug targets for diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  507 CTCAGTCTCAAAGAAAAAAGACAGGGAGCAAAGAGGACAAGAAAATACGTCTGG 560
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-490007/54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-FEB-2000;
21-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; G-protein subunit Gbeta3; variant; high blood pressure; asthma; hypertonia; cardiac infarction; coronary disease; heart disease; circulation disease; diabetes; psychiatric disease; Crohn's disease; immunological disease; psoriasis; colitis ulcerosa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SIFF/) SIFFERT W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-JUN-2000; 2000DE-1030945.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DE10030945-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transplant rejection; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTCTGTCTCAAAAAAAAAAAAAAAAAAAAAAAAAAAGACAGTAAGAAAGGGCTAG 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunological and psychiatric diseases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                    2000WO-EP00602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000DE-1004681.
2000DE-1007587.
99DE-1003489
                                                                                                                                                                                                                                                                           peptide; BNP; primer; human; detection; epitope;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 25.2;
Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               37 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein to including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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Вb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
 reflects relative abundance of corresp. mRNA in specific human
              Identifying gene signatures in 3'-directed human cDNA library for diagnosis of abnormal cell function, by preparing cDNA that
                                                                WPI; 1995-206931/27.
                                                                                            Matsubara K,
                                                                                                                       (ОКИВ/) ОКИВО К
                                                                                                                                                                                         11-NOV-1994;
                                                                                                                                                                                                                      01-JUN-1995.
                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                    cell typing;
                                                                                                                                   (MATS/) MATSUBARA K
                                                                                                                                                               12-NOV-1993;
                                                                                                                                                                                                                                                                                                          Gene signature; messenger RNA; mRNA; relative abundance; frequency; human; cloning; mapping; non-biased library; diagnosis; detection;
                                                                                                                                                                                                                                                                                                                                                           Human gene signature HUMGS03631.
                                                                                                                                                                                                                                                                                                                                                                                       02-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                    AAT22086;
                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT22086 standard; cDNA to mRNA; 89 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1838 TG 1839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1778 AGGGCGGCCTGCTGATGAAGCAGTACTACCAGTTCATGCTGCAGGAGGTTCTGGAGGACT 1837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                natriuretic peptide (BNP) in a sample using two antibodies, that recognize different epitopes of the N-terminal pro-BNP. The method can be used for diagnosing heart insufficiency. The recombinant N-terminal pro-BNP is useful as a standard in a method to detect N-terminal pro-BNP and also for production of antibodies against N-terminal pro-BNP. This sequence represents a primer used in the cloning of recombinant N-terminal pro-BNP which is described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 70 BP; 10 A; 19 C; 21 G; 20 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel method to detect N-terminal pro-brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Identification of N-terminal pro-brain natriuretic peptide (pro-BNP) using two antibodies that recognize different epitopes of the N-terminal pro-BNP, useful for diagnosis of heart insufficiency -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 32; 35pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HOFF ) ROCHE DIAGNOSTICS GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  68 TG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8 AGGGAGGTCTGTTCAACCTGCAGTTCGGACAGTTTACCCTGCAGGTGGTTACGCTGTTCC 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lill H, Stahl P,
                                                                                          Okubo K;
                                                                                                                                                                                                                                                                                                  abnormal cell function; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                               93JP-0355504
                                                                                                                                                                                        94WO-JP01916
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99DE-1011044.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.5%;
62.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Krueger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 25.2; DB 2: Pred. No. 4.9e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ~
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Borgya A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gallusser A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ٦1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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tissues

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RESULT 12
AAT92244
ID AAT92
    A single-stranded DNA (or its complementary strand or the corresp. double-stranded DNA) which comprises one of the 7837 "GS" sequences of your sequences of the 7837 "GS" sequences to part of the man genomic DNA, cDNA or mRNA is claimed. The GS (Gene Signature) sequences were obtained from 3'-directed CDNA ibraries prepared from various human tissues; synthesis of cDNA was initiated from the GS (Gene Signature) and of mRNA by using poly(T) as the sole primer. Since the 3'-C oriented coNAs hybridise with specific mRNAs. Each library C all the 3'-oriented coNAs hybridise with specific mRNAs. Each library is constructed so as to reflect accurately the relative abundance of different mRNAs in the particular tissue from which it was derived. The appearance frequency of a given GS in a cDNA library can be determined (esp. using primers and probes derived from the GS sequences) as a means of diagnosing abnormal cell function or for recognising different cell types.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 1039; 2245pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4545 GT 4546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 89 BP; 39 A; 8 C; 15 G; 27 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Metal ion resistance; mercuric ion reductase; mera; mercury; toxic heavy metal; transgenic plant; mutagenic; bacterial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mercuric ion reductase merA PCR mutagenesis primer 307-339S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-JAN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT92244 standard; DNA;
mutagenise the mera coding sequence derived from transposon {\tt Tn21}, adapt it for plant expressibility. The nucleic acid molecule pro
                                                                                                                                                                            WPI; 1997-470112/43.
                                                                                                                                                                                                               Meagher RB, Summers AO;
                                                                                                                                                                                                                                                     (UYGE-) UNIV GEORGIA RES FOUND INC
                                                                                                                                                                                                                                                                                          21-APR-1995;
                                                                                                                                                                                                                                                                                                                              21-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                   16-SEP-1997.
                                                                                                                                                                                                                                                                                                                                                                                                         US5668294-A
                                                                                                                                                                                                                                                                                                                                                                                                                                          Transposon Tn21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transposon; Tn21; PCR; ss.
                   The present sequence represents a PCR mutagenesis primer used to mutagenise the merA coding sequence derived from transposon Tn21.
                                                                                                                DNA encoding mercuric ion reductase in plant-expressible form producing transgenic plants resistant to toxic heavy metals
                                                                             Example 1; Column 43-44; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              73 GT 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                          95US-0427097
                                                                                                                                                                                                                                                                                                                                95US-0427097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 25.2;
Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5.8e+03;
ches 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 16; Length 89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
             produced
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 13
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        is useful for producing transgenic plants that are resistant to toxic heavy metals (especially mercury) and so can be used remediate and/or revegetate contaminated areas. The bacterial merA gene is derived from the transposon Tn21, which was originally isolated from the Incompatibility Group IncFII resistance plasmid NR1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 99 BP; 29 A; 21 C; 25 G; 24 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3026 ATGTCCTGACTAAATGTGAGAGTGGTAGAGGTACAAGACCCAGGTGGGCGACTCAAGGAT 3085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3086 TTGATTTTCTACAAGCAATTGAACCTGCCTTTAT 3119
                              The invention relates to a nucleic acid molecule comprising a coding sequence for an organometal resistance protein, which is operably linked downstream of and under the regulatory control of a plant-expressible transcription and translation regulatory sequence. The organometal resistance gene is especially the merA gene from the transposon Tm21 or the E.coll merB gene. merA encodes a mercuric ion reductase and merB encodes a organomercury lyase. The nucleic acid sequences are useful for producing transgenic plants which are capable of efficiently processing mercury. These plants are able to remove or detoxify metal compounds such as methyl mercury and ionic mercury from soil, sediment, and aquatic environments providing an efficient bioremediation of metal and organometal contaminated sites. This sequence represents a PCR primer used to amplify the Tm21 merA gene which is mutated by random mutagenesis (see AAZ28584-Z28589 for mutated soll and the mitated senior are avaraged in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Organometal; resistance; transcription; translation; regulation; mera; transposon; Tn2l; merB; mercuric ion reductase; organomercury lyase; ss; transgenic plant; detoxification; metal compound; soil; sediment; primer aquatic environment; bioremediation; contamination; PCR; amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Primer 307-339S for Tn21 merA gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-DEC-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ28594 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US5965796-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Transposon Tn21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                           Example 3; Column 17-18; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                   Metal resistance sequences for producing transgenic plants -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYGE-) UNIV GEORGIA RES FOUND INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-OCT-1999
              mutated sequences) and the mutated sequences are expressed in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           62 GTGAGCATACTCAAGCAAGCCAAGTTGCCTATAT 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 54.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 AAGACCCAGCTATAGGTGAAGCTGTTACTGCTGCATTTCGCATGGAAGGCATTGAAGTGC 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          51; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Meagher RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95US-0427097.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          97US-0878957.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Summers AO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 25.2; DB 18
Pred. No. 6.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 43; Indels 7 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 99;
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transgenic plants of the invention

В

Matches

0

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RESULT 14
AAA73946/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                      immediately upstream of a gfp coding sequence to form the synthetic gene. The amplified fragment was cloned into the mammalian expression vector pCDNA3, which contains SV40 orl and the CMV promoter, and was used in a The synthetic construct was introduced into COS-1 cells and expression of the reporter protein (green fluorescent protein) was measured. A series of 4 gfp reporter constructs was made in which the gfp gene is preceded entire set of isoaccepting codon triplets. Codons with a higher translational efficiency than their corresponding synonymous codons can be identified. These codons may then be used to replace the less protein undifferentiated epithelial cells such as COS-1 cells.
Sequence 54 BP; 9 A; 14 C; 20 G; 11 T; 0 other;
                                                                                                                                                                                                                               The present sequence is a primer used to generate a synthetic gfp gene by PCR amplification of a humanised gfp gene. A single artificial start codon followed by a stretch of five identical codons was fused in frame
                                                                                                                                                                                                                                                                                                   Example 1; Page 183; 190pp; English.
                                                                                                                                                                                                                                                                                                                             tandem repeats of the codon, and measuring expression -
                                                                                                                                                                                                                                                                                                                                        Determining translational efficiency of codons in cells, comprising introducing synthetic constructs with reporter genes fused in frame to
                                                                                                                                                                                                                                                                                                                                                                                                                            Zhou J, Frazer IH;
                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-499118/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JAN-2000; 2000WO-AU00008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Green fluorescent protein; GFP; reporter gene; codon utilisation; translational efficiency; protein abundance; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SUNX/) SUN X.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYQU ) UNIV QUEENSLAND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200042215-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GFP Leu(CTG)5 forward primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-DEC-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3026 ATGTCCTGACTAAATGTGAGAGTGGTAGAGGTACAAGACCCAGGTGGGCGACTCAAGGAT 3085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA73946;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA73946 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3086 TTGATTTTCTACAAGCAATTGAACCTGCCTTTAT 3119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 99 BP; 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        62 GTGAGCATACTCAAGCAAGCCAAGTTGCCTATAT 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAGACCCAGCTATAGGTGAAGCTGTTACTGCTGCATTTCGCATGGAAGGCATTGAAGTGC 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99AU-0008078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; 21 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  54 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25 G; 24 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 25.2;
Pred. No. 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6.2e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              43;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 99;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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Query Match Best Local Similarity

69.4%;

Pred. No. 4.7e+03 Score 25;

DB 21;

Length 54;

밁 Ϋ́ Вþ

ABL58788/c RESULT 16

0.5%;

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ABA69691/c
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                                                                                                                                                                            Matches
                                                                                                                                                                                                                       Query Match
                                          3875 AGCTC 3879
                                                                                                              3815 ACGAATTGAAAATATTCGAAGGCATCAAACACCCCCAATCTGGTTCGGTATTTTGGTGTGG 3874
                                                                                                                                                                                                                                                                         probe of the invention.

Note: The sequence data for this patent did not form part of printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a single exon nucleic acid probe for measuring human gene expression in a sample derived from human foetal liver. The single exon nucleic acid probes may be used for predicting, measuring and displaying gene expression in samples derived from human fetal liver. The present sequence is a single exon nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                    Sequence 91 BP; 28 A; 19 C; 13 G; 31 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; SEQ ID NO 17996; 639pp + sequence listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human fetal liver -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-483447/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-FEB-2000;
26-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JAN-2001; 2001WO-US00669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-AUG-2001
                                                                          67 AAGAAATTATTATGATGAAAGACTGTAAACACCCAAATATTGTTGCTTATTTTGGAAGCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200157277-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; foetal liver; gene expression; single exon nucleic acid probe; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human foetal liver single exon nucleic acid probe #17996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2732 CCAGTCAGCCGGTCATCGCCAAAGCTTTGCAGCAGCTGAAGAATGATGC 2780
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABA69691 standard; DNA; 91 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABA69691;
7 ATCTC
                                                                                                                                                                                            Match 0.5%;
Local Similarity 61.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    53 CCAGTGAACAGTTCCTCGCCCTTGCTCAGCAGCAGCAGCAGCATGGTAC
                                                                                                                                                                          40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hanzel DK,
                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000GB-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000US-0234687
2000US-0236359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000US-0180312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      200005-0632366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000US-0608408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000US-0207456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chen W,
                                                                                                                                                                      0;
                                                                                                                                                                                     Score 25; DB 22;
Pred. No. 6.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rank DR
                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                            DB 22; Length 91;
                                                                                                                                                                      25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15;
                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ٠.
                                                                                                                                                             0
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RESULT 17
AAC12392/c
ID AAC123
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cc using a microfluidics system (MS). A starter oligonucleotides (I) cc using a microfluidics system (MS). A starter oligonucleotide (ON) is cc eather bound (directly or through a spacer) to the surface of the cc reaction space in MS or is synthesised directly, with one end bound to cc the surface. ON corresponds to one end of (I) being synthesised. One or cc more rounds of hybridisation of 'sequemers', each corresponding to a cc segment of (I) or its complement and having overlapping ends that allow hybridisation, is performed. If required blunt ends, formed e.g. after a cpolymerase reaction, are converted to sticky ends suitable for cc hybridisation. Where there are nicks between consecutive sequemers, cc injation is performed or if a segment of one or more nucleotides (nt) is complementary to a segment of sequemer) this is completed by complementary to a segment of sequemer) this is completed by cc process is repeated as necessary to assemble (I). The method is used to produce synthetic genes or other bloodydically active nucleic acids. The cc process can be fully automated, under computer control. In an entire coligonucleotides (ABLS8783-ABLS8788) are used to synthetically generate the non-coding strand the Requoria victoria green fluorescence protein cc (GFP) complementary to the coding strand (ABLS6950).
                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic GFP oligonucleotide GFP-5b SEQ ID NO 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DE10046069-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GFP; green fluorescent protein; microfluidic system; synthetic gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABL58788 standard; DNA; 96 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthesis of polynucleotide in microfluidics system, useful for preparing synthetic genes, comprises stepwise addition of sequemers to immobilized starter oligonucleotide -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-SEP-2000; 2000DE-1046069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Aequoria victoria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-316770/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bernauer HS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BERN/) BERNAUER H S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-SEP-2000; 2000DE-1046069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; Page 13; 18pp; German.
                                                                                                                                                                                                                                                                       1352 GTAGACATTGTCTGACTTCTATTTATAGACCATTTGTAGACAAAGCACTGAAGCAGATGG 1411
                         AAC12392 standard; cDNA; 77 BP
                                                                                                                                                                                               1412 GGTTAAGAAAGTTAATTTTAA 1432
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 96 BP; 41 A; 12 C; 16 G; 27 T; 0 other;
                                                                                                                                                                                                                                             96 GTATACATTGTGTGAGTTATAGTTGTATTCCAATTTGTGTCCAAGAATGTTTCCATCTTC 37
                                                                                                                                                 36 TTTAAAATCAATACCTTTTAA 16
                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                            0.5%; sec. No. /c. 56.8%; pred. No. /c. Mismatches
                                                                                                                                                                                                                                                                                                                                                                        Score 25; DB 24; Length 96; Pred. No. 7e+03;
                                                                                                                                                                                                                                                                                                                                                       35; Indels
                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Qy
                                                                                                                                                                                                                                                                                                                                                                  Вb
                                                                                                                                                                                                                                                                                                                                                                                                             QY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    D
                                                                                                                                                                                                                                                              RESULT 18
AAS06930/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. No ORR has yet been conclusively identified within the present sequence. The 5' ESTs were prepared from identified within the present sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed CDNA ilbraries. Such ESTs are not well suited for isolating cDNA sequences and the 5' ends of mRNAs and even in those cases where longer cDNAs sequences have been obtained, the full 5' UTR is rarely included by ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and serverion and serverions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; 5' EST; expressed sequence tag; secreted protein; cDNA isolation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human secreted protein 5' EST, SEQ ID NO: 16467.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene therapy; chromosome mapping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-OCT-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC12392;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EP1033401-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID 16467; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and fo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-500381/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dumas Milne Edwards J, Duclert A, Giordano J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-FEB-2000; 2000EP-0200610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 77 BP; 27 A; 14 C; 15 G; 21 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression and secretion vectors.
                                                                                                                                                                                                                                          AAS06930 standard; DNA; 92 BP
                    nucleocapsid phosphoprotein; large polymerase; attenuated vaccine;
                                        Infectious chimeric parainfluenza virus; antigenic determinant;
                                                                                                                                               12-SEP-2001 (first entry)
                                                                                                                                                                                            AAS06930;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            301 GACTCAGTGGGTGGATCTTTGCCAAAAAATCAATTCCAGATGTGGATCTCAATAAGCCT 360
                                                                                        Reverse PCR primer for RSV A G gene insert.
                                                                                                                                                                                                                                                                                                                                                                                                                361 TACCTCAG 368
                                                                                                                                                                                                                                                                                                                                                                  14 TATCTCAG 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                              74 GACTGTGTAGGTGCATTTGTTTAAAAAAATATAACTCCCCCATTGCCTCTGGGATAAAGTCT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ocal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0122487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 24.8; DB 21;
Pred. No. 6.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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Length 77; Indels

0;

0

QΥ B QY

Db

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RESULT 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The Sequence is described in an invention relating to novel infectious chimeric parainfluenza viruses (PIVs). The virus comprises a major condimeric parainfluenza viruses (PIVs). The virus comprises a major condimeric parainfluenza viruses (PIVs). The virus comprises a major condimeric protein (I), an uncleocapsid phosphoprotein (P), a large genome, or antigenome combined with one or more heterologous genets) or genome, or antigenome combined with one or more heterologous genets) or compreheners of one or compreheners of the chimeric protein (P) is useful in an attenuated vaccine to elicit an immune compreheners of the chimeric protein (PIV may also elicit an immune composition compreheners of the chimeric protein (PIVs). The chimeric protein the first chimeric protein composition compos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KW KW KW XXX OSS OSS PN PN PN XXX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
gene therapy; chromosome mapping; ss
                                                                             Human secreted protein 5' EST, SEQ ID NO: 15574.
           Human; 5' EST; expressed sequence tag; secreted protein; cDNA isolation;
                                                                                                                                 06-OCT-2000 (first entry)
                                                                                                                                                                                                 AAC11499;
                                                                                                                                                                                                                                          AAC11499 standard; cDNA; 94 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 92 BP; 20 A; 12 C; 26 G; 34 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence for reverse PCR primer for respiratory syncytial virus (RSV) A G gene insert is used with the forward PCR primer (AAS06929) in the construction of bovine/humanPIV3-RSV chimeric viruses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isolated infectious chimeric parainfluenza virus (PIV), useful in an attenuated vaccine to elicits an immune response against one or more virus(es) selected from human PIVI (HPIVI), HPIV2 and HPIV3 -
                                                                                                                                                                                                                                                                                                                                                           45 TACCTCTCCCCCCAGCACACCTCGACAGATGAAACGCATGTCAACCAAACATCAGAGGAA 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 10; Page 150; 305pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-356173/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-DEC-1999;
10-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-DEC-2000; 2000WO-US33293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human respiratory syncytial virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-JUN-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human PIV1; HPIV1; HPIV2; HPIV3; RSV; pathogen; measles; PCR primer;
respiratory syncytial virus; respiratory tract infection; bovine; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9905-0459062
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       990S-0170195.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     63.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schmidt AC, Durbin AP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 24.8; DB 22; Pred. No. 7.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.7e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 92;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Skiadopoulos MH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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XX AC XX AC

21-FEB-2000; 2000EP-0200610

06-SEP-2000.

Q

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RESULT 20
AAC15541/c
                                                                                                                                                                                                                                                                                                   Db
                                                                                                                                                                                                                                                                                                                              QΥ
                                                                                                                                                                                                                                                                                                                                                             В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    XX OS XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                     Matchės
                                                                         Human; 5' EST; expressed sequence tag; secreted protein; cDNA isolation; gene therapy; chromosome mapping; ss.
                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                      1439 ACAAGCTAATGGATGGTTCCTTGCAAAGGGCA 1470
                                                                                                                                                                                                                                                                                                                                                                           1379 GACCATTTGTAGACAAGCACTGAAGCAGATGGGGTTAAGAAAGTTAATTTTAAGACTTC 1438
                                                                                                                           Human secreted protein 5' EST, SEQ ID NO: 19616
                                                                                                                                                          06-OCT-2000 (first entry)
                                                                                                                                                                                            AAC15541;
                                                                                                                                                                                                                  AAC15541 standard; cDNA; 94 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mRNAS encoding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences cDNA sequences have been obtained, the full 5' UTR is rarely included. USETS are derived from mRNAs with intact 5' ends and can therefore be in diagnostic, forensic, gene therapy and chromosome mapping procedures. Extraction upstream regulatory sequences and to design
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 94 BP; 46 A; 9 C; 14 G; 25 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression and secretion vectors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence is one of a large number of 5' ESTs derived from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID 15574; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-500381/45.
                                                                                                                                                                                                                                                                                          62 AAGATTTAATGTAGGAAATATTATTAAAGGCA 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dumas Milne Edwards J, Duclert A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-FEB-2000; 2000EP-0200610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-SEP-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                               2 GAAAATATAAAAATACTAACCAAAAGAAGTCTGAGGTAACTATGGTAATATTAAAACTCC
                                                                                                                                                                                                                                                                                                                                                                                                                  50;
                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99US-0122487
                                                                                                                                                                                                                                                                                                                                                                                                                           0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                     Score 24.8; DB 21,
                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Giordano J;
                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                             42;
                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 94;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             71'
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RESULT 21
ABN54107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and servertion vectors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; SEQ ID 19616; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1595 TGTCGTGGGAGGAGCTGAAGGCCATGGATTTACCTTCATTCGAACCTGCCTTCCTAGTTC 1654
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and fo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-500381/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dumas Milne Edwards J, Duclert A, Giordano J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1655 TCTGCC 1660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 94 BP; 35 A; 10 C; 29 G; 16 T; 4 other;
                                                                                                                                                                                                                                                                                                                                                                   15-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                      ABN54107;
                                                                                                                                                                                                                                                                                                                                                                                                                                       ABN54107 standard; DNA; 65 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression and secretion vectors.
                                                                                                                                                                                                            WO200210449-A2.
                                                                                                                                                                                                                                                                           Human; mouse; rat; splice transcript; detection; RNA transcript; splice variant; transcriptome; oligonucleotide library; ss.
                                                                                                                                                                                                                                                                                                                                Mouse spliced transcript detection oligonucleotide SEQ ID NO:26855.
              Shoshan A, Wasserman A, Mintz E,
                                                                                 28-JUL-2000; 2000US-221607P
02-MAY-2001; 2001US-287724P
                                                                                                                                                                           07-FEB-2002.
                                                                                                                                                                                                                                             Mus musculus
                                                   (COMP-) COMPUGEN INC
                                                                                                                                        20-JUL-2001; 2001WO-IB01903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10 CCTTCC 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  70 TTTGGTGATTCAAGTTGAGTGCATTACTTTTCCCTTCCTKCCTKCCTTCCTTCCTTCCTHCTTK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0122487.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 24.8; DB 21;
Pred. No. 7.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                  Mintz L,
                    Faigler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 94;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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The present invention describes oligonucleotide libraries for detecting cc messenger RNAs that populate a (sub-)transcriptome, where the cc (sub-)transcriptome comprises messenger RNAs transcribed from multiple transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to a set of messenger RNAs transcribed from a given transcription unit of the poligonucleotide libraries are useful for detecting mRNAs from a cc biological sample, in expression profiling studies, in qualitatively or quantitatively characterising the corresponding transcriptome, and in cc detecting RNA transcripts and splice variants of human or animal cc transcriptomes. The libraries may also be used as specialised mini cc particular biological or pathological state, and so allowing the corresponding transcriptome under a cc particular biological or pathological state, and so allowing the corresponding transcriptome under a cc detection of tissue—and pathology-specific genes such as those genes conly expressed in specific tissue under a specific pathological condition; to detect developmental specific genes such as those genes confictions and splice variants of a transcriptome of a patient suffering cranscripts and splice variants of a transcriptome of a patient suffering cranscripts and splice variants of a transcriptome of a patient suffering cc ondivided to the present invention.

CC N.B. The sequence data for this patent did not form part of the printed cat fip.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT65208/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription unit of a genome, useful for detecting tissue-, pathology-, and
20-MAR-1996;
02-JUN-1995;
02-JUN-1995;
05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-257383/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; SEQ ID 26855; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 developmental-specific genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2001 AGAAATAACTCATTACATACGGGGAGGAGGAAGCACAGGCCGGGAAGCTTTTCTGT 2055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 65 BP; 22 A; 9 C; 21 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                    identification; SELEX; anti-mitogenic; inhibition; cell; Systematic Evolution of Ligands by Exponential enrichment; epithelial; proliferation; diagnosis; treatment; fibroids;
                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                   epirnelial; proliferation; diagnosis; treatment; fibroids; kidney; lung; liver; dermal scarring; restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                       Transforming; growth factor; beta-1; TGF-beta-1; binding ligand;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Transforming growth factor beta-1 binding ligand D 11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT65208 standard; DNA; 71 BP
                                                                                                                          30-MAY-1996;
                                                                                                                                                                      05-DEC-1996
                                                                                                                                                                                                                     WO9638579-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT65208;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
     96US-0618693.
95US-0458423.
95US-0458424.
95US-0465591.
                                                                                                                       96WO-US08014
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 24.6; DB pred. No. 7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 65;
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Best Local
                P-PSDB; AAB01926
                              WPI; 2000-338602/29
                                                          Young MW,
                                                                                    (UYRQ ) UNIV ROCKEFELLER
                                                                                                                   19-JUN-1998;
                                                                                                                                              19-JUN-1998;
                                                                                                                                                                             02-MAY-2000
                                                                                                                                                                                                                                                                      ATP binding domain; Doubletime; DBT; casein kinase family protein; serine/threonine kinase; Period; PER; phosphorylation; stability; circadian rhythm; clock; sleep disorder; depression; narcolepsy;
                                                                                                                                                                                                                                   Drosophila melanogaster
                                                                                                                                                                                                                                                                                                                                 cDNA encoding Drosophila Doubletime (DBT) ATP binding domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3958 GAACATGTGAGTTAGGCTGTATTCAAAGCAGCAGCACCATTGCGATCAACGTCCTCC 4012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the remainder of the mixture. The partitioned nucleic acids were then amplified to yield a mixture of nucleic acids enriched for sequences with higher affinity and specificity for binding to TGF-beta-1. The ligand is anti-mitogenic and may be used to inhibit epithelial cell proliferation, or in the diagnosis and treatment of TGF-beta-1 mediated pathological conditions, e.g. fibrotic conditions such as fibroids of the kidney, lung and liver and more
                                                                                                                                                                                                                                                                                                                                                                    18-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                           AAA52344 standard; cDNA; 72 BP
                                                                                                                                                                                                                                                                   jet lag; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 71 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of Ligands by Exponential enrichment (SELEX). Briefly a candidate mixture of nucleic acids was contacted with TGF-beta-1, and nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence, a transforming growth factor beta-1 (TGF beta-1) binding ligand, was identified by Systematic Evolution
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           acute conditions such as dermal scarring and restenosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           acids having an increased affinity to TGF-beta-1 partitioned from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 15; Page 123; 209pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identification of nucleic acid ligands to TGF-beta, PDGF and hKGF using SELEX, used in the diagnosis and treatment of proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1997-034387/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (NEXS-) NEXSTAR PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             56 GAAGACGTAAGTAAGTATTAAAAACATACCCCCCCTGGCCGCATCGTCCTCC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ocal Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                36;
                                                        Blau J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Janjic N,
                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                 98US-010066
                                                                                                                                             98US-0100664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15 A; 16 C; 22 G; 18 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95US-0479783.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                95US-0465594.
95US-0479725.
                                                        Price J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pagratis N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 24.6; DB 18
Pred. No. 7.5e+03;
                                                     Kloss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ringquist S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Toothman PJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC (AMB01927). It plays a role in regulating the concentration of the clock CP protein Period (PER). PER acts together with Timeless (TIM) in the CP protein Period (PER). PER acts together with Timeless (TIM) in the CP protein Period (PER). PER acts together with Timeless (TIM) in the CP protein of the circadian rhythm. The PER protein is normally unstable in the cytoplasm, but is stabilised on heterodimerisation with TIM. The CP per and time stabilised on heterodimerisation with TIM. The CP per and time per and time genes through a negative feedback loop. Because PER/TIM heterodimers are only present in large numbers when high CP levels of per and time RNA have accumulated, self-sustained oscillations are produced in the negative feedback loop. DBT plays a key role in the CP several sites, with a high level of PER monomers. It phosphorylates PER at CP several sites, with a high level of PER phosphorylation correlating CP with increased instability and degradation. The invention also CP associated with a lengthened (slower) circadian rhythm. The DBT-L (AABO1924 and CP associated with a lengthened (slower) circadian rhythm. The DBT-S mutant CP associated with a lengthened (slower) circadian rhythm. The DBT-S mutant CP appears to phosphorylate PER more rapidly than the wild-type protein, CP associated with a lengthened (slower) circadian rhythm. DBT agonists of DBT-L causes hypophosphorylation and persistence of monomeric PER, while the action CP contains rhythm disorders including depression, narcolepsy, jet lag, and other sleep-wake ovele mental disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           matches 42; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                       Trypanosoma; Toxoplasma; Giardia;
                                                                                                                                                                                                                              viral infection; human immunodeficieny virus; metanoma; bacterial infection; Salmonella; Legionella; parasitic infection;
                                                                                                                                                                                                                                                                                Savine; vaccine; cancer; viral infection; HIV; hepatitis C virus;
                                                                                                                                                                                                                                                                                                                                  HCV DNA encoding HepCla segment 133.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3754 ATGGCCATGAA 3764
                                                                                                                                                                                                                                                                                                                                                                                       08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                  ABK36568;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABK36568 standard; DNA; 90 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3694 ATTGGAGAAGGCCAGTATGGGAAGGTGTACACCTGCATCAGCGTCGACACCGGGGAGCTG 3753
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 72 BP; 18 A; 19 C; 23 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            other sleep-wake cycle mental disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence represents cDNA encoding the ATP binding domain of the Drosophila circadian rhythm protein Doubletime (DBT). DBT (AAB01923) is a member of the casein kinase protein family and comprises an ATP binding domain (AAB01926) and a serine/threonine kinase catalytic domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 25; Column
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel nucleic acids encoding clock protein, doubletime which is useful for treating disorders of circadian rhythm such as jet lag and mental disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 GTGGCCATCAA 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 ATAGGATCGGGATCGTTCGGCGACATCTACCTGGGCACCACGATCAACACTGGCGAGGAG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  71-72; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 24.6; DB 21
Pred. No. 7.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 72;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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26-MAY-2000; 2000AU-0007761 25-MAY-2001; 2001WO-AU00622. WO200190197-A1 Hepatitis C virus

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cc rhe invention relates to a new synthetic polypeptide (1) comprising cc several different segments of at least one parent polypeptide linked cc together in a different relationship relative to their linkage in the cc parent polypeptide to impede, abrogate or otherwise alter at least one cc function associated with the parent polypeptide and for inducing an cc immune response against a pathogen or cancer. Also included are a cc synthetic polypucteotide encoding and a computer system for cd designing the synthetic polypeptides. The synthetic polypeptides and cc golypucleotides are referred to as a Savine. The synthetic polypeptide is cc useful for modulating immune responses preferably directed against a perfect of the lung, breat, ovary, cervix, cc colon, head and neck, pancreas, prostate, stomach, bladder, kidney, bone cc immunodeficiency virus), hepatitis, influenza, Japanese encephalitis cc virus, Epstein Barr virus and respiratory syncytial virus) bacterial cc e.g., infections caused by Neisseria, Meningococcal, Haemophilus, cc salmonella, Streptococcal, Legionella and Mycobacterium or parasitic crypanosoma, Toxoplasma and Giardia) infections. The present cc construct a savine of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                               Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Дb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Qy
Вþ
                                                                                                                                                                                                                                                                                                                                     RESULT 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Fig 26; 364pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New synthetic polypeptides having several different segments of at least one parent polypeptide linked together differently compared to the linkage in the parent polypeptide, for inducing immune response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Thomson SA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (AUSU ) UNIV AUSTRALIAN NAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      against a pathogen or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  P-PSDB;
                                                                                                      Neutrophil; gene expression profile; granulocyte; pathogen-exposed; sterile inflammatory disease; detection; therapeutic agent; human; expression modulator; pathogenic infection; cell activation; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                 2334 GAAATCAAAAC 2344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2274 CCTGGAAATAGCAGCAGAATTCAGGCTTTCAGCCCCAGTTAGAGACCTCCTGGATGTTCT 2333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 90 BP; 22 A; 23 C; 21 G; 24 T; 0 other;
                   WO9910536-A1
                                                                                                                                                                                     Human neutrophil cDNA clone 876.
                                                                                                                                                                                                                          17-JUN-1999 (first entry)
                                                                                                                                                                                                                                                                                                  AAX23469 standard; DNA; 98 BP
                                                      Homo sapiens
                                                                                       global transcriptional response; ss
                                                                                                                                                                                                                                                                                                                                                                                              20 CANATCCANTC 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        80 CCAGGCAGTTGGGGCATCAGTTTGGCTTTCAGCCAGGTTTTGAAATCGCTGAGCACTTCC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2002-147575/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAU84730
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ramshaw IA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.5%; Score 24.6; DB 24; 59.2%; Pred. No. 8.8e+03;

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 90;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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$\text{SQ} \text{ $\text{CQ} \text{ $\text{ $\text{CQ} \text{ $\text{ $\text{ $\text{CQ} \text{ $\text{ 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 26
AAX25128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  QΥ
                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a method for the comparison of gene expression profiles from granulocytes from a test subject and from pathogen-exposed or sterile inflammatory disease granulocytes or quiescent granulocytes. The method is used to detect pathogen exposure or sterile inflammatory disease in a subject and to identify therapeutic agents that modulate expression of a gene in response to a pathogenic infection or to sterile inflammatory disease in a subject. The method tests for global transcriptional response of granulocytes during cell activation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-MAR-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4971 GGTTTATTTTA 4981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4911 GTTTGTAATGCAAAAGGCTGATTACTGAAATTTAAGAAAAGGTTCTTTTTTCAATAAAT 4970
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 98 BP; 30 A; 16 C; 20 G; 32 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Page 41; 84pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detection of pathogen exposure or sterile inflammatory disease in a subject - by comparing gene expression profiles from granulocytes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-204678/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Goguen J, Newburger P, Prashar Y, Weissman SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYYA ) UNIV YALE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of the patient and control granulocytes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnosis; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Flavivirus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        80%E Bundle peptide-encoding oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX25128;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX25128 standard; DNA; 82 BP
(HAWA-) HAWAII BIOTECHNOLOGY GROUP INC
                                                              31-JUL-1997;
                                                                                                                                27-JUL-1998;
                                                                                                                                                                                               11-FEB-1999
                                                                                                                                                                                                                                                              WO9906068-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                 misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              33 GTTACAATATA 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93 GTTTATCATACATAAGACCGCATGACTATAATACAAAGGGGGGGTTCTTTTTTCATTTAAC 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.5%;
Local Similarity 59.2%;
es 42; Conservativo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             dengue virus; envelope protein; vaccine; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97US-0056844
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98WO-US17284.
                                                                     97US-0904227
                                                                                                                                    98WO-US15447.
                                                                                                                                                                                                                                                                                                                                                          /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                       /*tag=
                                                                                                                                                                                                                                                                                                                        "single stranded overhang on complementary strand of sequence 5'-TCGA-3'"
                                                                                                                                                                                                                                                                                                                                                                                   .
מ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 24.6; DB 20
Pred. No. 9.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Yerramilli SV;
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RESULT 27
ABA68615/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                         Human genome-derived single exon nucleic acid probes useful analyzing gene expression in human fetal liver -
                                                                    WPI; 2001-483447/52
                                                                                                                                                                                     03-AUG-2000;
                                                                                                                      (MOLE-)
                                                                                                                                                             21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                  26-MAY-2000;
30-JUN-2000;
                                                                                                                                                                                                                           04-FEB-2000;
                                                                                                                                                                                                                                                    30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         construction of an expression vector for secretion of non-covalently linked 80%E Bundle, where 80%E comprises a C-terminally truncated envelope (E) protein (see AAY05522) of dengue virus serotype 2 strain PR159/S1, and Bundle is the linker and helix-turn-helix moiety that is designed to form a non-covalently associated 4-helix bundle domain that enhances the dimerisation of the 80%E molecule. A vaccine for protecting against DEN-2 infection comprises a dimeric 80%E protein that has been secreted as a recombinant protein from a eukaryotic cell.
                                                                                                                                                                                                                                                                                                       WO200157277-A2
                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                                                                                                                                                                      01-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1789 CTGATGAAGCAGTACTACCAGTTCATGCTGCAGGAGGTTCTGGAGGACTTCGGAGGAAGC 1846
                                                                                                                                                                                                                                                                                                                                                                                                                                   ABA68615;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABA68615 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 82 BP; 19 A; 19 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This double-stranded oligonucleotide was annealed to another double-stranded oligonucleotide (see AAX25127) to generate a Esequence encoding the C-terminal half of a linker and helix-turn-helix domain (see AAY05526). This was utilised in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Recombinant dimeric flaviviral envelope vaccine - c dimeric 80%E protein, useful for protecting against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 4; Page 30; 60pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          especially dengue virus infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Coller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 CTGCTGAAGGGCCCCCGCAAGGGCGAGCTGGAGGAGCTGCTGAAGCACCTGAAGGAGC
                                                                                                                                                                                                                                                                                                                                                                         foetal liver single exon nucleic acid probe #16920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1999-153454/13
                                                                                                                                                                                                                                                                                                                                                       toetal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           37;
                                                                                                                     MOLECULAR DYNAMICS INC
SEQ ID NO 16920; 639pp + sequence listing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                              Hanzel
                                                                                                                                                           2000US-0180312.
2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0632366.
2000US-0234687.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                              2000GB-0024263.
                                                                                                                                                                                                                                                  2001WO-US00669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Harada KE,
                                                                                             DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                  gene expression; single exon nucleic acid probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.5%;
                                                                                           Chen W, Rank DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IVY JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       32 G; 12 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 24.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   McDonnell M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peters
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        generate a DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
 English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   flavivirus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              82;
                                        for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               XXCCCCCCXX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                   26-MAY-2000;
30-JUN-2000;
30-AUG-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
                                                                                      Claim 4; SEQ ID No 14054; 530pp; English.
                                                                                                                             Single exon nucleic acid probes for analyzing gene
                                                                                                                                                                                      Penn SG,
                                                                                                                                                                                                               (MOLE-)
                                                                                                                                                                                                                                                                                                                      04-FEB-2000;
                                                                                                                                                                                                                                                                                                                                              30-JAN-2001; 2001WO-US00666
                                                                                                                                                                                                                                                                                                                                                                         09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     congenital heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; gene expression; heart; microarray; vascular syste cardiovascular disease; hypertension; cardiac arrhythmia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Probe #14054 for gene expression analysis in human heart cell sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABA35588 standard; DNA; 99
                                                                                                                                                                                                                                                                                                                                                                                                  WO200157274-A2
                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABA35588;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a single exon nucleic acid probe for measuring human gene expression in a sample derived from human fliver. The single exon nucleic acid probes may be used for predimensuring and displaying gene expression in samples derived from fetal liver. The present sequence is a single exon nucleic acid probe of the invention.

Note: The sequence data for this patent did not form part of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format direction wipo at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            751 AATGGTACTTCAGTAGAAGGGCAGTGCAAAGCCACTCCTGGAACAAAGATTGTAGGTTAC 810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              811 TCAACACATCATGA 824
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                           2001-488899/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TGACCTCATCATCA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AATATGAATGCCGTAGCATGGGATCCCAAGCCCTGCCCACGACCAATGCATGTACCTCTC
                                                                                                                                                                                                              MOLECULAR DYNAMICS
                                                                                                                                                                                   Hanzel DK,
                                                                                                                                                                                                                                     2000US-0180312.
2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0234685.
2000US-0236359.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 25 A; 14 C; 35 G; 25 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.5%;
58.1%;
                                                                                                                                                                                   Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                     88
                                                                                                                                                                                   ž
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 24.4; DB 22;
Pred. No. 1.1e+04;
0; Mismatches 31;
                                                                                                                                                                                   Rank DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           microarray; vascular system;
                                                                                                                             expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 format directly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 predicting,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     from human
                                                                                                                             ij
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            foetal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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The present invention relates to single exon nucleic acid probes for measuring human gene expression in a sample derived from human heart. The present sequence is one such probe. The probes may be used for present sequence is one such probe.

predicting, measuring and displaying gene expression in samples derived from the human heart via microarrays. By measuring gene expression, the

diagnosing,

probes are useful for predicting,

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RESULT 29
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QΥ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                 Matches
                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               monitoring and prognosing diseases of the human heart and vascular system e.g. cardiovascular disease, hypertension, cardiac arrhythmias and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 99 BP; 25 A; 14 C; 35 G; 25 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      congenital heart disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human insulin derivative; diabetes; Zinc ion complex GlyA21 AspB3 PCR primer #127; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human insulin derivative GlyA21 AspB3 PCR primer #127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-NOV-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ86411;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ86411 standard; DNA; 100 BP
   Sequence 100 BP; 29 A; 23 C; 22 G; 26 T; 0 other;
                                                                                                                       The two subunits of the derivative are covalently connected via disulphide bonds between Cys residues A7/B7 and A20/B19. The derivative, which may be present as a zinc ion complex, can be used as a fast action treatment for diabetes.
                                                                                                                                                                                   AAQ86399 and AAQ86411 are a pair of primers for the PCR amplification of the Asp3 B-chain/Gly21 A-chain human insulin derivative DNA.
                                                                                                                                                                                                                                                              Acylated insulin deriv. which may be present as a Zinc ion complex - is used to treat diabetes and is rapid acting.
                                                                                                                                                                                                                                                                                                                                                                                                                     02-FEB-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9507931-A
                                                                                                                                                                                                                                  Example 3; Page 34; 100pp; English.
                                                                                                                                                                                                                                                                                                           WPI; 1995-131314/17.
                                                                                                                                                                                                                                                                                                                                           Markussen
                                                                                                                                                                                                                                                                                                                                                         Andersen AS,
                                                                                                                                                                                                                                                                                                                                                                                    (NOVO ) NOVO-NORDISK AS.
                                                                                                                                                                                                                                                                                                                                                                                                                                    17-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      811 TCAACACATCATGA 824
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   751 AATGGTACTTCAGTAGAAGGGCAGTGCAAAGCCACTCCTGGAACAAGATTGTAGGTTAC 810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          34 TGACCTCATCATCA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94 AATATGAATGCCGTAGCATGGGATCCCAAGCCCTGCCCACGACCAATGCATGTACCTCTC 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match 0.5%; focal Similarity 58.1%;
                                              ocal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 43;
                                 49; Conservative
                                              h 0.5%; Score 24.4; DB 16;
Similarity 54.4%; Pred. NO. 1.1e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                         Halstrom JB, Havelund S,
                                                                                                                                                                                                                                                                                                                                                                                                                     93DK-0001044
94US-0190829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94WO-DK00347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 24.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .1e+04;
                                                                                                                                                                                                                                                                                                                                                            Jonassen I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31;
                                   41; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 99;
                                                                 Length 100;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                     0,
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                                     Gaps
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6 A 6

48

RESULT 31 AAK21231/c

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABA72801/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 30
δÃ
                                                  밁
                                                                                                                                                                                 Query Match
                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4529 TATTTACATAAAGACTGTGCTGAGAAGCAG 4558
                                                                                                                                                                                                                                                                   The invention relates to a single exon nucleic acid probe for measuring human gene expression in a sample derived from human foetal liver. The single exon nucleic acid probes may be used for predicting, measuring and displaying gene expression in samples derived from human fetal liver. The present sequence is a single exon nucleic acid probe of the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABA72801;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABA72801 standard; DNA; 80 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        W0200157277-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; foetal liver; gene expression; single exon nucleic acid probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human foetal liver single exon nucleic acid probe #21106
                                                                                       3462 AAGAAGAAGTTGGGAACTTCGGACACTAATCAGCCAGAGTAAAGATACTGCTTCTAAACT 3521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; SEQ ID NO 21106; 639pp + sequence listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human fetal liver -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-483447/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-SEP-2000; 2000US-0236359.
04-OCT-2000; 2000GB-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JAN-2001; 2001WO-US00669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
  3522 A 3522
                                                                                                                                                                                                                              Sequence 80 BP; 18 A; 15 C; 13 G; 34 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Penn SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   37 CAATTGGAAAACTACTGTGGTTAGACGCAG 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97 AAGTCTGACGACGCTAAGGGTATCGTTGAACAATGTTGTACTTCTATCTGTTCTTTGTAC 38
                                             Local Similarity
                                                                                                                                            38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hanzel DK,
                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000US-0207456
2000US-0608408
2000US-0632366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000US-0180312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000US-0234687
                                                                                                                                                            0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chen W, Rank DR;
                                                                                                                                              0;
                                                                                                                                                              Score 24.2; DB 22;
Pred. No. 1.1e+04;
                                                                                                                                              Mismatches
                                                                                                                                                 23;
                                                                                                                                                                                           Length 80;
                                                                                                                                                 Indels
                                                                                                                                                 0
                                                                                                                                                 Gaps
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AAK47388/c
ID AAK473
        XX EXX DXXX
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                                                                                                                                                                                                                                                                                                            RESULT 32
                                                                                                                                                                                                                                     QΨ
                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                Matches
                       Human bone marrow expressed single exon probe SEQ ID NO: 21945.
        Human;
                                                       06-NOV-2001
                                                                                                   AAK47388 standard; DNA; 80
                                                                                                                                                                                       3522 A 3522
                                                                                                                                                                                                                      3462 AAGAAGAAGTTGGGAACTTCGGACACTAATCAGCCAGAGTAAAGATACTGCTTCTAAACT 3521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
                                                                                                                                                                                                                                                                                                                                     probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia, epilepsy and cancers. The present sequence is one of the probes of the
                                                                                                                                                                                                                                                                                                          Sequence 80 BP; 18 A; 15 C;
                                                                                                                                                                                                                                                                                                                                                                                                  The present invention provides a number of single exon nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Single exon nucleic acid probes for analyzing gene expression in human
                                                                                                                                                                                                                                                                                                                                                                                                                                Example 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          brains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-483446/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Penn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001; 2001WO-US00667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200157275-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       epilepsy; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  microarray; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; brain expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human brain expressed single exon probe SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAK21231 standard; DNA; 80
                                                                                                                                                                                                        66 AAGAAATTATCAAGCACTGTGGACATTTAAAGGCCACAGAAATCATCCTGGTTATAAATT 7
                                                                                                                                                               6 A 6
                                                                                                                                                                                                                                                                       Local Similarity
     bone marrow expressed
                                                                                                                                                                                                                                                            38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULAR DYNAMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO: 21222; 650pp + Sequence Listing; English
                                                                                                                                                                                                                                                           Conservative
                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000US-0180312.
2000US-0207456.
2000US-068408.
2000US-0632366.
2000US-0234687.
2000US-02346359.
2000US-0236359.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                     0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chen W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           exon; gene expression analysis;
                                                                                                                                                                                                                                                                                                       13 G; 34 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
     exon;
                                                                                                                                                                                                                                                       Score 24.2; DB 22;
Pred. No. 1.1e+04;
0; Mismatches 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rank DR;
gene expression analysis; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                multiple sclerosis; schizophrenia;
                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21222
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RESULT 33
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                                                                                                                                                                                                                                             AAI53226/c
                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
04-FEB-2000;
                     30-JAN-2001; 2001WO-US00663.
                                                                                                                                                  Probe #21912 used to measure gene expression in human placenta sample
                                                 09-AUG-2001.
                                                                                                                      genetic
                                                                                                                             Probe; microarray; human; placenta; antenatal diagnosis,
                                                                      WO200157272-A2
                                                                                                                                                                               17-OCT-2001
                                                                                                                                                                                                         AAI53226;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention provides a number of single exon nucleic acid probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow samples, which may enable the improved diagnosis and treatment of cancers such as lymphoma, leukaemia and myeloma. The present sequence is one of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
                                                                                                                                                                                                                               AAI53226 standard; DNA;
                                                                                                                                                                                                                                                                                                                3522 A 3522
                                                                                                                                                                                                                                                                                                                                                        3462 AAGAAGAAGTTGGGAACTTCGGACACTAATCAGCCAGAGTAAAGATACTGCTTCTAAACT 3521
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 80 BP; 18 A; 15 C; 13 G; 34 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            the probes of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   analyzing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 4; SEQ ID NO: 21945; 658pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human bone {\tt marrow} -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JAN-2001; 2001WO-US00668
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200157276-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     microarray; cancer; leukaemia; lymphoma; myeloma;
                                                                                                                                                                                                                                                                                                                                   66 AAGAAATTATCAAGCACTGTGGACATTTAAAGGCCCACAGAAATCATCCTGGTTATAAATT
                                                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                           6 A 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001-488900/53.
                                                                                                                                                                                                                                                                                                                                                                                       38;
                                                                                                                      disorder;
                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; 2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0234685.
2000US-0236359.
2000US-0024263.
                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2000US-0180312
                                                                                                                                                                                                                                                                                                                                                                                                0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chen W,
                                                                                                                                                                                                                              80 BP
                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                               Score 24.2; DB 22
Pred. No. 1.1e+04;
                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rank DR;
                                                                                                                                                                                                                                                                                                                                                                                                            DB 22;
                                                                                                                                                                                                                                                                                                                                                                                     23;
                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                            80;
                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                 0;
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RESULT 34
AAA922
ID AAA92
XX AAA92
XX Thyme
XX Thyme
XX Thyme
XX Unid
XX OS-J
PX 30-L
XX 30-L
XX WPI
PR WPI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-MAY-2000; 2000US-0207456.
30-JUN-2000; 2000US-0658408.
30-JUN-2000; 2000US-052366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0236359.
04-OCT-2000; 2000GB-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MOLE-) MOLECULAR DYNAMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to single exon nucleic acid probes (SENP). The present sequence is one such probe. The probes are useful for producing a microarray for predicting, measuring and displaying gene producing a microarray for predicting, measuring and displaying gene expression in samples derived from human placenta. The probes are useful expression in samples derived from human genetic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-488897/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Penn SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 25; SEQ ID No 21912; 654pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 80 BP; 18 A; 15 C; 13 G; 34 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3462 PAGAAGAAGTTGGGAAGTTCGGACACTAATCAGCCAGAGTAAAGATACTGCTTCTAAACT 3521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3522 A 3522
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA92397 standard; DNA; 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Thymosin alpha-1 nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-JAN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA92397;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rhymosin alpha-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CN1258747-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   66 AAGAARTATCAAGCACTGTGGACATTTAAAGGCCACAGAAATCATCCTGGTTATAAATT 7
                                                                                                                                                                                                                                                                                                                       30-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                          30-DEC-1998;
                                                                                                                                                                                                                                                                       (UYFO-) UNIV NO 4 MILITARY MEDICAL
                                                                                                                                                  WPI; 2000-558962/52
P-PSDB; AAB23792.
                                                                                                                                                                                                                          Zhao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6 A 6
                                               Claim 1; Page 2; 9pp; Chinese.
                                                                                            Preparation of 4-tandem thymosin alpha-1 with colon bacillus
The present invention describes the preparation of 4-tandem thymosin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  genome-derived single exon nucleic acid probes useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.5%;
Similarity 62.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene expression in human placenta
                                                                                                                                                                                                                                Zhang Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                98CN-0113062
                                                                                                                                                                                                                                                                                                                                                                                98CN-0113062
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          T-alpha-1; 4-tandem thymosin alpha-1; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chen W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 24.2; DB 2;
Pred. No. 1.1e+04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rank DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 80;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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Дb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    alpha-1 (T-alpha-1) gene synthesising process comprising cloning the gene sequence, measuring and other steps. The T-alpha-1 genes are gene sequence, measuring and other steps. The T-alpha-1 genes are connected serially into tetramer and expressed with colon bacillus. Through ion exchange purification, biological active T-alpha-1 matter with high purity and high activity is obtained. The present invention with high purity and high activity is obtained. The present at the inestimable market demand. The present sequence represents the inestimable market demand. The present sequence.
      3718 GTGTACACCTGCATCAGCGTCGACACCGGGGAGCTGATGGCCATGAAAGAGAT 3770
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 84 BP; 26 A; 21 C; 32 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABK36388 standard; DNA; 90 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV DNA encoding GAP A segment 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-MAY-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABK36388;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Savine; vaccine; cancer; viral infection; HIV; hepatitis C virus; viral infection; human immunodeficieny virus; melanoma; viral infection; Salmonella; Legionella; parasitic infection; bacterial infection; Salmonella; ds. Trypanosoma; Toxoplasma; Giardia; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus type 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Locar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200190197-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        35
The invention relates to a new synthetic polypeptide (I) comprising several different segments of at least one parent polypeptide linked together in a different relationship relative to their linkage in the stogether in a different relationship relative to their linkage in the parent polypeptide to impede, abrogate or otherwise alter at least one function associated with the parent polypeptide and for inducing an function associated with the parent polypeptide and for included are a synthetic polynucleotide encoding and a computer system for synthetic polypeptides. The synthetic polypeptides and solupton the synthetic polypeptide is designing the synthetic polypeptide is as a savine. The synthetic polypeptide is polynucleotides are referred to as a savine. The synthetic golypeptide is polypeptide in mountained in munumer partners of the lung, breast, ovary, cervix, useful for modulating immune responses preferably directed against a polynucleotides. The synthetic polypeptide is designed in the synthetic polypeptide is polypeptide. The synthetic polypeptide is polypeptide is a savine. The synthetic polypeptide is polypeptide is a savine of the lung, breast, ovary, cervix, useful for modulating immune responses preferably directed against a polypeptide is polypeptide. The synthetic polypeptide is the lung, breast, ovary, cervix, useful for modulating immune responses preferably directed against a polypeptide. The synthetic polypeptide is designed in the synthetic polypeptide in the synthetic polypeptide is designed in the syntheti
                                                                                                                                                                                                                                                                                                                                                                                                                                             New synthetic polypeptides having several different segments of at least one parent polypeptide linked together differently compared to the linkage in the parent polypeptide, for inducing immune response against a pathogen or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-MAY-2000; 2000AU-0007761
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAY-2001; 2001WO-AU00622.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (AUSU ) UNIV AUSTRALIAN NAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P-PSDB; AAU84549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thomson SA,
                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Fig 12; 364pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002-147575/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ramshaw IA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.5%;

 Mismatches

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Pred. No. 1.1e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 84;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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Дδ
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          AAH51601 represents a human genomic nucleotide sequence comprising sbg1, 934665, sbg2, g35017 and g35018 nucleic acid sequences located on the human chromosome 13q31-q33 locus. The nucleotide sequences contain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Qy
biallelic markers and polymorphisms. Sequences AAH51602 - AAH51626 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8888888888<del>%</del>&
                                                                   Claim 1; Page 563; 737pp; English.
                                                                                         markers are used for genotyping and detecting schizophrenia or bipolar
                                                                                      disorder and predisposition to these disorders
                                                                                                                                             WPI; 2000-619082/59.
                                                                                                                                                                                                                                                                                        30-MAR-1999;
30-APR-1999;
30-APR-1999;
                                                                                                                                                                                                          (GEST ) GENSET.
                                                                                                                                                                                                                                            29-JUL-1999;
29-JUL-1999;
                                                                                                                                                                                                                                                                                                                               30-MAR-2000; 2000WO-IB00435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                  27-JUL-1999;
                                                                                                                                                                                                                                                                              14-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                              05-OCT-2000.
                                                                                                                                                                                                                                                                                                                                                                                  WO200058510-A2
                                                                                                                                                                                                                                                                                                                                                                                                        Pan troglodytes.
                                                                                                                                                                                                                                                                                                                                                                                                          biallelic marker; polymorphism; schizophrenia; bipolar disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sbg1 exon sequence SEQ ID 57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-AUG-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH51645;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH51645 standard; DNA; 94 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4100 GTTCAGTAAAGCTCAAAAACAATGCCCAGACCATGCCTGGTGAAGTGAACAGCACCCTGG 4159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4160 GGA 4162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Compositions comprising the polypeptide may be used in the treatment or prophylaxis against viral (such as infections caused by HIV (human virus, Epstein-Barr virus), hepatilis, influenza, Japanese encephalitis (e.g., infections caused by Neisseria, Meningococcal, Haemophilus, Salmonella, Streptococcal, Legionella and Mycobacterium or parasitic (e.g., infections caused by Plasmodium, Schistosoma, Leishmania, Trypanosoma, Toxoplasma and Giardia) infections. The present construct a savine of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 90 BP; 29 A; 19 C; 15 G; 19 T; 8 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             83 WGA 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23 GGTCCGAGAATYTCACARACAATGYCAAAACCATTATCGTCCAMCTCAACRAAAGCGTCG 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Socal Similarity
                                                                                                                                                                                                                                                                                                                                                                                                          934665; sbg2; g35017; g35018; chromosome 13q31-q33; haplotype;
                                                                                                                                                                              Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       35,
                                                                                                                                                                                                                         99US-0145915.
99US-0146452.
99US-0146453.
99US-0162288.
                                                                                                                                                                                                                                                                       99US-0132065.
99US-0143928.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                 99US-0126903
99US-0131971
                                                                                                                                                               Chumakov I, Bougueleret L, Bihain B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           55.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           // Score 24.2; DI
// Pred. No. 1.2e-
5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.2e+04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 90;
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RESULT 37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polynucleotides comprising sequences from sbg1 and g35018 biallelic markers are used for genotyping and detecting schizophrenia or bipol disorder and predisposition to these disorders -
                                                                                WPI; 2000-619082/59.
                                                                                                                                                                                                                                                                       30-MAR-1999;
30-APR-1999;
30-APR-1999;
                                                                                                                                   Cohen D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                              (GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                               29-JUL-1999;
                                                                                                                                                                                                                              29-JUL-1999;
                                                                                                                                                                                                                                            27-JUL-1999;
                                                                                                                                                                                                                                                           14-JUL-1999;
                                                                                                                                                                                                                                                                                                                            30-MAR-2000; 2000WO-IB00435.
                                                                                                                                                                                                                                                                                                                                                                                              W0200058510-A2
                                                                                                                                                                                                                                                                                                                                                                                                                              sbg1; g34665; sbg2; g35017; g35018; chromosome 13q31-q33; haplotype;
biallelic marker; polymorphism; schizophrenia; bipolar disorder; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                           Gorilla gorilla.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sbg1 exon sequence SEQ ID 80.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-AUG-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAH51668;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH51668 standard; DNA; 94 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1048 AAGAATTTATCAGACATTGGCTGGCCAGTGTTTGAAA 1084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 94 BP; 29 A; 15 C; 20 G; 30 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         46 AAGAATTTCTCAGATATTTGCTGGGCACTTTATCGAA 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                           Blumenfeld м,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                       99US-0162288.
                                                                                                                                                                                                        99US-0146453
                                                                                                                                                                                                                                     99US-0143928
99US-0145915
                                                                                                                                                                                                                      99US-0146452
                                                                                                                                                                                                                                                                 99us-0132065
                                                                                                                                                                                                                                                                                              9908-0126903.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.5%;
78.4%;
                                                                                                                      Chumakov I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 24.2; DB 21;
                                                                                                               Bougueleret L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.2e+04
                                                                                                           Bihain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 94;
               or bipolar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
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A AAHS1601 represents a human genomic nucleotide sequence comprising sbyl, cc g34665, sbg2, g35017 and g35018 nucleic acid sequences located on the CC g34665, sbg2, g35017 and g35018 nucleic acid sequences contain cc human chromosome 13g31-g33 locus. The nucleotide sequences contain cc biallelic markers and polymorphisms. Sequences AAHS1602 - AAHS1626 and cc AAB62907 - AAB62915 represent cDNA human sbyl cDNA sequences and protein products. Primers AAHS1632 - AAHS1639 are used cc cDNA sequences and protein products. Primers AAHS1632 - AAHS1699 are used cc cDNA sequences AHHS1642 - AAHS1693 - AAHS1693 are used cc amplicons which comprise biallelic markers located on the chromosome cc amplicons which comprise biallelic markers located on the chromosome cc are represented in the sequences by degenerate/undefined base codes. PCR cc 13g31-g33 locus are represented in AHS1819 are used in the isolation of sequences of cc are represented in AAHS1819 are used in the isolation of sequences of cc invention. The biallelic marker containing nucleotide sequences are cc for genotyping by determining the identity of the nucleotide at a biallelic marker in cc contained by a sequence. The nucleotide are biallelic marker in a biological sample from single or multiple cc subjects. By determining the frequency of a biallelic marker in a cc contained to determine a predisoposition to or early onset of schozophrenia or cc contain against schizophrenia or bipolar disorder or a beneficial response to or side effects related to cc freatment against schizophrenia or bipolar disorder.
RESULT 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAH51692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 568; 737pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1048 AAGAATTTATCAGACATTGGCTGGCCAGTGTTTGAAA 1084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 94 BP; 29 A; 15 C; 20 G; 30 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sbgl; g34665; sbg2; g35017; g35018; chromosome 13q31-q33; haplotype;
biallelic marker; polymorphism; schizophrenia; bipolar disorder; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sbg1 exon sequence SEQ ID 104.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-AUG-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH51692 standard; DNA; 94 BP
                                                                                                                                                                                                                                                                                                                                                                                             WO200058510-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                     Macaca mulatta
                                                                                                                                                                                                                                                                                                           30-MAR-2000; 2000WO-IB00435.
                                                                                                                                                                                                                                                                                                                                                     05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-MAR-1999;
30-APR-1999;
                      Cohen D,
Essioux L;
                                                                                                                                                                                                                                30-APR-1999;
                                                                                  (GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OCAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                               Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                99US-0126903.
99US-0131971.
99US-0132065.
99US-0143928.
99US-0145915.
99US-0146452.
99US-0146453.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.5%;
                                                    Chumakov I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 24.2;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred
                                                      Bougueleret L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .2e+04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 94;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                              Bihain B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
```

В QУ

Matches

Local 29;

Conservative

0; Mismatches

Indels

0;

1048 AAGAATTTATCAGACATTGGCTGGCCAGTGTTTGAAA 1084

46 AAGAATTTCTCAGATATTTGCTGGGCACTTTATGGAA 82

δÃ

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XX AAH51601 represents a human genomic nucleotide sequence comprising sbg1, cg 34665, sbg2, g35017 and g35018 nucleic acid sequences located on the cg g34665, sbg2, g35017 and g35018 nucleic acid sequences contain chuman chromosome 13q31-q33 locus. The nucleotide sequences contain challelic markers and polymorphisms. Sequences AAH51602 - AAH51602 and protein products. AAH51627 - AAH51631 and AAB62915 - AAB62915 represented protein products. Primers AAH51632 - AAH51628 are used cc changlicons which comprise biallelic markers located on the chromosome camplicons which comprise biallelic markers located on the chromosome camplicons which comprise biallelic markers located on the chromosome camplicons which comprise biallelic markers located on the chromosome cc anglicons which comprise biallelic marker located on the chromosome cc are represented in the sequences by degenerate/undefined base codes. PCR cromosome comprise biallelic marker containing nucleotide sequences of cc used to determine the identity of the nucleotide at a biallelic marker in cc used to determine the identity of the nucleotide at a biallelic marker in cc subjects. By determining the identity of a nucleotide at a Region cc of primary products. By determining the identity of a nucleotide at a Region cc of primary and a predisoposition to or early onset of schizophrenia or cc used to determine a predisoposition to or early onset of schizophrenia or cc used to determine a predisoposition to or early onset of schizophrenia or cc created biallelic marker in a biological sample effects related to cc a trait and a phenotype and a trait can be detected. The sequences can be created to determine a predisoposition to or early onset of schizophrenia or cc created to determine a predisoposition to or early onset of schizophrenia or cc created to determine a predisoposition to or early onset of schizophrenia or cc containing the identity of the sequences can be contained to determine a predisoposition to or early onset of schizophrenia or cc treatment agai
                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polynucleotides comprising sequences from sbg1 and g35018 biallelic markers are used for genotyping and detecting schizophrenia or bipolar disorder and predisposition to these disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 573; 737pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-619082/59.
                                                                             Sequence 94 BP; 28 A; 16 C; 20 G; 30 T; 0 other;
Similarity
       0.5%;
       Score 24.2;
Pred. No. 1
                  .2e+04;
                                                DB 21;
                                                Length 94;
```

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RESULT 39
ABA71915/c
ID ABA71915 standard; DNA; 95 BP.
                                   04-FEB-2000; 2000US-0180312.

26-MAY-2000; 2000US-0204456.

30-JUN-2000; 2000US-0608408.

31-JUN-2000; 2000US-0632366.

21-SEP-2000; 2000US-0234687.

21-SEP-2000; 2000US-0236359.
                                                                                                                                                                                                                                       Human; foetal liver; gene expression; single exon nucleic acid probe;
                                                                                                                                                                                                                                                                 Human foetal liver single exon nucleic acid probe #20220.
                                                                                                                                                                                                                                                                                             01-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                       ABA71915;
                                                                                                                                                            09-AUG-2001.
                                                                                                                                                                                     WO200157277-A2
                                                                                                                                                                                                                Homo sapiens
                                                                                                                                30-JAN-2001; 2001WO-US00669.
                          04-OCT-2000; 2000GB-0024263
(MOLE-) MOLECULAR DYNAMICS INC
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AAK20302/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Qγ
                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DЬ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Дb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-FEB-2000; 2000US-0180312
26-MAY-2000; 2000US-0207456
30-JUN-2000; 2000US-0608408
03-AUG-2000; 2000US-062366
21-SEP-2000; 2000US-0234687
27-SEP-2000; 2000US-0236359,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
Single exon nucleic acid probes for analyzing gene expression in human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                     WPI; 2001-483446/52.
                                                                 Penn SG,
                                                                               (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                           04-OCT-2000;
                                                                                                                                                                                                     30-JAN-2001; 2001WO-US00667
                                                                                                                                                                                                                                                                                                 Human; brain expressed exon; gene expression analysis; probe; microarray; Alzheimer's disease; multiple sclerosis; schizophrenia;
                                                                                                                                                                                                                                                       WO200157275-A2
                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                    05-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                        Human brain expressed single exon probe SEQ ID NO: 20293.
                                                                                                                                                                                                                                                                                                                                                                                                                 AAK20302 standard; DNA; 95 BP
                                                                                                                                                                                                                                                                                                                                                                                                  AAK20302;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1202 ATCCTAGGG 1210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1142 AAGGAGAATTAAAGGAGTTGGAAAGTAGTACGGATGAAGAGGAAGAAGGAACAAATCTCTG 1201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 95 BP; 21 A; 28 C; 13 G; 33 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  printed specification, but was obtained in electronic
from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probe of the invention.

Note: The sequence data for this patent did not form part of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a single exon nucleic acid probe for measuring human gene expression in a sample derived from human foetal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               measuring and displaying gene expression in samples derived from human fetal liver. The present sequence is a single exon nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; SEQ ID NO 20220; 639pp + sequence listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human fetal liver -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Penn SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28 AGGCTAAGG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-483447/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             41;
                                                            Hanzel DK,
                                                                                                      2000GB-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                        Chen W, Rank DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chen W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 24.2; DB 22;
Pred. No. 1.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rank DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 95;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              format directly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Job time :
     Search completed: November 11, 2002, 07:16:44
                                                                                                   Qγ
                                                                    В
                                                                                                                                      DЬ
                                                                                                                                                                    Qγ
                                                                                                                                                                                                                                                                                      XX CCCCCCCX XX XX XX
                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                     Query Match
                                                                                                                                             1142 AAGGAGAATTAAAGGAGTTGGAAAGTAGTACGGATGAGAGTGAAGAAGAAGAACAAATCTCTG 1201
                                                                                              1202 ATCCTAGGG 1210
                                                                                                                                                                                                                                                      Sequence 95 BP; 21 A; 28 C; 13 G; 33 T; 0 other;
                                                                                                                                                                                                                                                                                                              The present invention provides a number of single exon nucleic acid probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia,
                                                                                                                                                                                                                                                                                               epilepsy and cancers. The present sequence is one of the probes of the
                                                                                                                                                                                                                                                                                                                                                                                                                   Example 4; SEQ ID NO: 20293; 650pp + Sequence Listing; English.
                                                                                                              88 ACGGAGAAGTAATGGAATATAAGACTGGACAGCAAGGACGTGATTATGAAGAAATGTGTC 29
                                                               28 AGGCTAAGG 20
                                                                                                                                                                                                                  Local Similarity
660 secs
                                                                                                                                                                                                   41;
                                                                                                                                                                                               Conservative
                                                                                                                                                                                                          0.5%;
                                                                                                                                                                                        0;
                                                                                                                                                                                                       Score 24.2; DB 22;
Pred. No. 1.2e+04;
                                                                                                                                                                                        Mismatches
                                                                                                                                                                                        28;
                                                                                                                                                                                      Indels
                                                                                                                                                                                                                     Length 95;
                                                                                                                                                                                     0
                                                                                                                                                                               Gaps
```